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Minimum DB
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Perfect score:
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| (cgn2 6/ptodata/1/pubpna/US07 PUBCOMB.seq:*
| (cgn2 6/ptodata/1/pubpna/PCT NEW PUB.seq:*
| (cgn2 6/ptodata/1/pubpna/US06 NEW PUB.seq:*
| (cgn2 6/ptodata/1/pubpna/US06 PUBCOMB.seq:*
| (cgn2 6/ptodata/1/pubpna/US07 NEW PUB.seq:*
| (cgn2 6/ptodata/1/pubpna/PCTUS PUBCOMB.seq:*
| (cgn2 6/ptodata/1/pubpna/PCTUS PUBCOMB.seq:*
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Copyright (c) 1993 - 2004 Compugen Ltd.
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S US-10-024-444B-1
S US-10-017-161-635
US-09-886-055-198
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US-10-343-650A-513
S US-10-343-650A-513
S US-10-30-846-25
US-09-974-591-11
US-09-974-591-13
US-09-777-789-10
US-09-974-149-11
US-09-974-149-11
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                                Sequence 1, Appli
Sequence 635, App
Sequence 198, App
Sequence 198, App
Sequence 198, App
Sequence 39, Appl
Sequence 35, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 10, Appl
Sequence 11, Appl
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                      Sequence
Sequence
      Sequence
      8, A<sub>1</sub>
                    App1:
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RESULT 1 US-10-024-444B-1

GENERAL INFORMATION:
APPLICANT: Padigaru, Muralidhara
APPLICANT: Gerlach, Valerie L.
APPLICANT: Gerlach, 71ennda Sequence 1, Application US/100244448 Publication No. US20030165858A1

Smithson,

Stone,

David Glennda

Conley, Tart, Matthew Bin-Yang, Ruey Conley, Pamela B.

Tomlinson, James E

ALIGNMENTS

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US-10-024-444B-1
                                                                                                                                                   SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                 APPLICANT
 Best Local Similarity 100. Matches 977; Conservative
                                  Query Match
                                                                                                                                                                                                                                                                         APPLICANT: Topper, James N.
APPLICANT: Kekuda, Ramesh
APPLICANT: Casman, Stacie J.
APPLICANT: MacDougall, John R.
APPLICANT: Shlomit, Edinger R.
TITLE OF INVENTION: No. US20030165858A1el GPCR-Like Proteins and Nucleic Acids Encod
TITLE OF INVENTION: Same
FILE REFERENCE: 21402-224 AG
                                                                                           LENGTH: 977
TYPE: DNA
ORGANISM: Human
                                                                                                                                                                NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver. 2
                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/024,444B
CURRENT FILING DATE: 2002-12-19
PRIOR APPLICATION NUMBER: 60/256635
PRIOR FILING DATE: 2000-12-18
100.0%; Score 977;
100.0%; Pred. No. 0;
tive 0; Mismatches
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   Mismatches
                                        DB 15;
   0;
                                      Length 977;
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; Sequence 635, Application US/10017161; Publication No. US20030143668A1
                       RESULT 2
US-10-017-161-635
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APPLICANT: SUWA, MAXIKO
APPLICANT: ASAI, KIYOSHI
APPLICANT: ASAI, KIYOSHI
APPLICANT: ASAI, KIYOSHI
APPLICANT: ASURATANI, HIROYUKI
ITITLE OF INVENTION: NOVEL & PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 084335/0152
CURRENT APPLICATION NUMBER: US/10/017,161
CURRENT FILING DATE: 2002-12-18
PRIOR APPLICATION NUMBER: UP 2001/246789
PRIOR FILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 2430
SOFTWARE: PATENTIN VEY: 2.1
SEQ ID NO 635
LENGTH: 1033
TYPE: DNA
ORGANISM: Homo sapiens
FRATURE:
NAME/KEY: SQUICE
LOCATION: (1)..(1033)
FRATURE:
COCATION: (201)..(833)
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Similarity 99.9%;
                                                         TACGTGACAGGTGTGACTTTCCTCTTGCTCCCCATTTCTGCCATTGTGGCCTCCTACACA
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                       ACCTGCTCTTCCCACCTGATTGTGGTCGGGATGTTCTATGGAGCTGCCACACTTCATGTAT
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TITLE OF INVENTION: HUMAN OLFACTORY RECEPTORS AND GENES ENCODING SAME
FILE REFERENCE: P 0278005
CURRENT APPLICATION NUMBER: US/09/804,291
CURRENT FILING DATE: 2001-03-13
PRIOR FILING DATE: 2000-03-13
PRIOR FILING DATE: 2000-03-24
PRIOR PILING DATE: 2000-03-24
PRIOR PILING DATE: 2000-04-14
PRIOR PILING DATE: 2000-04-14
PRIOR PILING DATE: 2000-04-14
PRIOR PILING DATE: 2000-04-16
PRIOR PILING DATE: 2000-04-17
PRIOR PILING DATE: 2000-05-26
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-886-055-198
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US-09-886-055-198
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APPLICANT: ZOZULYA, SERGEY
APPLICANT: ZOZULYA, SERGEY
TITLE OF INVENTION: RECEPTOR FINGERPRINTING, SENSORY PERCEPTION, AND
TITLE OF INVENTION: BIOSENSORS OF CHEMICAL SENSANTS
FILE REFERENCE: 078003-0277150
CURRENT APPLICATION NUMBER: US/09/886,055
CURRENT APPLICATION NUMBER: US/09/886,055
CURRENT FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: 60/213,812
PRIOR TILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 522
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 198
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 198, Application US/09804291
Publication No. US20030088059A1
GENERAL INFORMATION:
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GENERAL INFORMATION:
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PRIOR PRIOR
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                                                                                                                                                                                                                                                                                                                                                                                         ZOZULA, SERGEY
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RESULT 6
US-10-220-382-39
JS-10-220-382-39
Sequence 39, Application US/10220382
Publication No. US20030119111A1
GENERAL INFORMATION:
APPLICANT: INCTE GENOMICS, INC.
APPLICANT: LAL, Preeti

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RESULT 5
US-10-343-650A-513
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PRIOR APPLICATION NUMBER: 60/226,534
PRIOR FILING DATE: 2000-08-16
PRIOR APPLICATION NUMBER: 60/230,732
PRIOR FILING DATE: 2000-09-07
PRIOR APPLICATION NUMBER: 60/266,862
PRIOR FILING DATE: 2001-02-07
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SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 198
LENGTH: 951
                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 513
LENGTH: 951
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Tocal Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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Matches 1
                                                                                                                                                    Matches
                                                                                                                                                                      Query Match
Best Local :
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PRIOR FILING DATE: 2001-02-13
NUMBER OF SEQ ID NOS: 694
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: HAGA, TATSUVA
TITLE OF INVENTION: NOVEL G-PROTEIN COUPLED RECEPTOR
FILE REFERENCE: 31671-186347
CURRENT APPLICATION NUMBER: US/10/343,650A
CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION NUMBER: JP 2000/237818
PRIOR FILING DATE: 2000-08-04
PRIOR FILING DATE: 2000-08-04
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                    886 CTACAGCCTGAGGAATAAGGAGGTCATGCGGGGCCTTGAGGAGGAGGGTCCTGGGAAAATACAT 945
                                                                                              826 ACAAGACAACATCATCTCTGTTTTCTACACAATTGTCACTCCAGCCCTGAATCCACTCAT 885
                                                                          807 ACAAGACAACATCATCTCTGTTTTCTACACAATTGTCACTCCAGCCCTGAATCCACTCAT
                                                                                                                                                    al Similarity 100.0%; I 120; Conservative 0;
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CTACAGCCTGAGGAATAAGGAGGTCATGCGGGCCTTGAGGAGGGTCCTGGGAAAATACAT
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                                                                                                                                                                    12.3%; Score 120; DB 13; 100.0%; Pred. No. 2.2e-51;
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Pred. No. 2.2e-51;
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PATTERSON, Chandra

Monique G.

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; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 25
; LENGTH: 951
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-300-846-25
                                                                                                                          APPLICANT: HAN, YI
APPLICANT: ECCULYA, SERGEY
APPLICANT: ZOZULYA, SERGEY
APPLICANT: ZOZULYA, SERGEY
APPLICANT: WANG, KUN
TITLE OF INVENTION: OLFACTORY RECEPTORS FOR ISOVALERIC ACID AND RELATED
TITLE OF INVENTION: MALODORANTS AND USE THEREOF IN ASSAYS FOR
TITLE OF INVENTION: IDENTIFICATION OF BLOCKERS OF MALODOR
TITLE OF INVENTION NUMBER: US/10/300,846
CURRENT APPLICATION NUMBER: US/10/300,846
CURRENT FILING DATE: 2002-01-16
PRIOR APPLICATION NUMBER: 00/348,371
PRIOR APPLICATION NUMBER: 00/809,291
PRIOR APPLICATION NUMBER: 00/809,291
PRIOR APPLICATION NUMBER: 60/341,872
PRIOR FILING DATE: 2001-03-13
PRIOR FILING DATE: 2001-12-21
NUMBER: 05/341,872
PRIOR FILING DATE: 2001-12-21
NUMBER: 05/341,872
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; Sequence 25, Application US/10300846

; Publication No. US20030207337A1
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; NAME/KEY: misc feature
; OTHER INFORMATION: Incyte ID No. US20030119111A1 7472446CB1
US-10-220-382-39
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Best Local Similarity
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SOFTWARE: PERL Program
SEQ ID NO 39
LENGTH: 951
                                                                                                                   PERIOR FILING DATE: 2001-12-21
NUMBER OF SEQ ID NOS: 86
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ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TRIBOULEY, Catherine
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100.0%; Pred. No. 2.2e-51;
vative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 15; Length 951;
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; LOCATION: (27)..(998)
US-09-974-591-11
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Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local Similarity
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                                     APPLICANT: Alsobrook II, John P
APPLICANT: Burges, Catherine E
APPLICANT: Grosse, William M
APPLICANT: Lepley, Denise M
APPLICANT: Padigaru, Muralidhara
APPLICANT: Spytek, Kimberly A
APPLICANT: Sytek, Kimberly A
APPLICANT: Sytek, Kimberly A
TITLE OF INVENTION: NO. US20030059830Alel Single Nucleotide Polymorphisms for Olfactor
TITLE OF INVENTION: Receptor-like Polypeptides and Nucleic Acids Encoding
TITLE OF INVENTION: the Same
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APPLICANT: Burgess, Catherine E
APPLICANT: Grosse, William M
APPLICANT: Lepley, Denise M
APPLICANT: Padigaru, Muralidhara
APPLICANT: Padigaru, Muralidhara
APPLICANT: Syrtek, Kimberly A
APPLICANT: Syrtek, No. US20030059830Alel Single Nucleotide Polymorphisms for Olfactor
TITLE OF INVENTION: Receptor-like Polypeptides and Nucleic Acids Encoding
TITLE OF INVENTION: the Same
FILLE REFERENCE: 15966-654 CIP
CURRENT APPLICATION NUMBER: US/09/974,591
CURRENT FILING DATE: 2001-10-09
PRIOR APPLICATION NUMBER: 60/245,292
PRIOR FILING DATE: 2000-11-02
NUMBER OF SEQ ID NOS: 26
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Best Local Similarity
CURRENT APPLICATION NUMBER: US/09/974,591
                         FILE REFERENCE: 1
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                         .5966-654 CIP
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US-09-974-149-11
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                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                  Sequence 11, Application US/09974149 Publication No. US20030175705A1
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SEQ ID NO 10
LENGTH: 1008
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Best Local Similarity
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   APPLICANT:
APPLICANT:
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                                       APPLICANT: Alsobrook II, John P
APPLICANT: Burgess, Catherine I
APPLICANT: Grosse, William M
APPLICANT: Lepley, Denise M
APPLICANT: Padigaru, Muralidhar
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TITLE OF INVENTION: NOVEL POLYPEPTIDES AND NUCLEIC ACIDS ENCODING SAME FILE REFERENCE: 15966-654

CURRENT APPLICATION NUMBER: US/09/777,789

CURRENT FILING DATE: 2001-05-11
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PRIOR APPLICATION NUMBER: 60/180,646
PRIOR FILING DATE: 2000-02-07
NUMBER OF SEQ ID NOS: 101
CONTURBER.
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NUMBER OF SEQ ID NOS: 26
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Burgess, Catherine E
Grosse, William M
Lepley, Denise M
Padigaru, Muralidhara
Spytek, Kimberly A
Bader, Joel S
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                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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; LOCATION: (27)..(998)
US-09-974-149-13
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US-09-974-149-13
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APPLICANT: Alsobrook II, John P
APPLICANT: Burgess, Catherine E
APPLICANT: Grosse, William M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13, Application US/09974149 Publication No. US20030175705A1
                                                                                                                                                                                                                                     Matches
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Best Local Similarity
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Best Local Similarity 100.0%;
Matches 120; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Padigaru, Muralidhara
APPLICANT: Spytek, Kimberly A
APPLICANT: Bader, Joel S
APPLICANT: Baneal, Aruna
TITLE OF INVENTION: Methods of Use for No. US20030175705A1el Single Nucleotide
TITLE OF INVENTION: Polymorphisms of Olfactory Receptor-like Polypeptides
TITLE OF INVENTION: and Nucleic Acids Encoding the Same
FILE REFERENCE: 15966-654UB
CURRENT APPLICATION NUMBER: US/09/974,149
CURRENT FILING DATE: 2002-05-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/323,755
PRIOR FILING DATE: 2001-09-20
NUMBER OF SEQ ID NOS: 26
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TYPE: DNA
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LOCATION: (27)..(998)
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ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                     120;
CTACAGCCTGAGGAATAAGGAGGTCATGCGGGCCTTGAGGAGGGTCCTGGGAAAATACAT 945
                                                                                                                               ACAAGACAACATCATCTGTTTTCTACACAATTGTCACTCCAGCCCTGAATCCACTCAT
                                                                                  ACAAGACAACATCATCTGTTTTCTACACAATTGTCACTCCAGCCCTGAATCCACTCAT
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-777-789-8
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                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: JP 2001/246789
PRIOR FILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 2430
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 8, Application US/09777789
Publication No. US20030087815A1
GEMERAL INFORMATION:
                                                                                              Matches 120;
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Best Local Similarity
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                                                                                                                                      Query Match
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NAME/KEY: SOUTCE
LOCATION: (1)...(1351)
FEATURE:
NAME/KEY: CDS
LOCATION: (201)...(1151)
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PRIOR APPLICATION NUMBER: 05-11
PRIOR APPLICATION NUMBER: 60/180,646
PRIOR FILING DATE: 2000-02-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/017,161
CURRENT FILING DATE: 2002-12-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: SUWA, MAKIKO
APPLICANT: ASAI, KIYOSHI
APPLICANT: AKIYAMA, YUTAKA
APPLICANT: ABURATANI, HIROYUKI
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 084335/0152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 101
SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                           LENGTH: 1351
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                  Local Similarity
           1007 ACAAGACAACATCATCTGTTTTCTACACAATTGTCACTCCAGCCCTGAATCCACTCAT 1066
                              826 ACAAGACAACATCATCTCTGTTTTCTACACAATTGTCACTCCAGCCCTGAATCCACTCAT 885
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                                                                                              Conservative
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rematches 0;
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US-10-292-798-275
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APPLICANT: SUWA, MAKIKO
APPLICANT: ASIA, KIYOSHI
APPLICANT: AKIYAMA, YUTAKA
APPLICANT: AKIYAMA, YUTAKA
APPLICANT: ABURATANI, HIROYUKI
TITLE OF INVENTION: GUNAOSINE TRIDHOSPHATE-BINDING PROTEIN COUPLED RECEPTORS
FILE REFERENCE: 084335/166
CURRENT APPLICATION NUMBER: US/10/292,798
CURRENT FILING DATE: 2002-11-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 275
LENGTH: 1351
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                                                                                                                                                                                                                                               Matches 120;
                                                                                                                                                                                                                                                                                      Query Match
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PRIOR FILING DATE: 2001-12-18
PRIOR APPLICATION NUMBER: JP 2001-246789
PRIOR FILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 2070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE:
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NAME/KEY: CDS
LOCATION: (201)..(1151)
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LOCATION: (1)..(1351)
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Local Similarity 100.0%; P
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5. US20030235833A1
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Pred. No. 2.1e-51;
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Maximum DB seq length: 200000000
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Perfect score:
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            Published Applications AA:*

1: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

5: /cgn2_6/ptodata/2/pubpaa/PCTUS_PUBCOMB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/PCTUS_PUBCOMB.pep:*

7: /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

8: /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

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16: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

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  14 US-10-024-444B-2
14 US-10-017-161-636
19 US-09-886-055-197
10 US-09-894-291-197
10 US-09-912-976-53
1 US-09-965-422-8
2 US-10-343-650A-514
4 US-10-220-382-18
4 US-10-220-382-18
5 US-10-017-161-310
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Sequence 2, Appli
Sequence 636, App
Sequence 197, App
Sequence 9, Appli
Sequence 53, Appli
Sequence 53, Appli
Sequence 54, Appli
Sequence 514, App
Sequence 10, Appli
Sequence 310, Appli
Sequence 26, Appli
Sequence 276, Appli
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10	10	10	9	14	15	11	11	10	11	10	15	14	11	11	10	10	10	10	10	14	10	14	14	11	11	11	10	10	10
US-09-907-218-4	US-09-907-218-2	US-09-804-291-201	US-09-886-055-201	US-10-023-597-68	US-10-005-041A-105	US-09-981-566A-71	US-09-965-422-59	US-09-795-271-49	9-965-	-271-	US-10-005-041A-104	US-10-024-444B-7	US-09-965-422-56	-976	US-09-795-271-48	US-09-777-789-47	US-09-777-789-46	US-09-777-789-42	US-09-777-789-41	US-10-024-444B-4	US-09-795-271-51	US-10-024-444B-6	US-10-024-444B-5	US-09-965-422-60	US-09-912-976-51	US-09-912-976-50	US-09-974-149-14	US-09-974-149-12	US-09-777-789-11
4.	2	201	201,	68, 7	105	e 71,	59,	49,	e 57,	Sequence 50, Appl	104	7,	56, App	52, App	Sequence 48, Appl	47,	46,		41,	4,	51,	ص	5	60,	51,	50,	14,	Sequence 12, Appl	-

ALIGNMENTS

US-10-024-444B-2

Sequence 2, Application US/10024444B Publication No. US20030165858A1

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Tomlinson, James E.

--LICANT: Topper, James M.

APPLICANT: Kekuda, Ramesh
APPLICANT: MacDougall, John R.

APPLICANT: MacDougall, John R.

APPLICANT: MacDougall, John R.

APPLICANT: Shlomit, Edinger R.

ITITLE OF INVENTION: No. US20030165858A1el GPCR-Like Proteins and Nucleic Acids Encod

TITLE OF INVENTION NUMBER: US/10/024,444B

CURRENT APPLICATION NUMBER: US/10/024,444B

CURRENT FILING DATE: 2002-12-19

PRIOR APPLICATION NUMBER: US/10/024,444B

CURRENT FILING DATE: 2000-12-18

NUMBER OF SEQ ID NOS: 12

SOPTWARE: Patentin Ver. 2.1

EEQ ID NO 2

LENGTH: 316

TYPE: D^^
                                                                          US-10-024-444B-2
Query Match
Best Local Similarity 100.
Matches 316; Conservative
                   100.0%; Score 316; DB 14; 100.0%; Pred. No. 2.6e-284;
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•
     Mismatches
   0
•
                                      Length 316;
     Indels
     0;
   Gaps
       0
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0; Indels Length 316

0;

Gaps

0

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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: SUMA, MAKIKO
APPLICANT: ASAI, KIYOSHI
APPLICANT: ASAI, KIYOSHI
APPLICANT: ABURATANI, HIROYUKI
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 084335/0152
CURRENT APPLICATION NUMBER: US/10/017,161
CURRENT FILING DATE: 2002-12-18
PRIOR APPLICATION NUMBER: JP 2001/246789
PRIOR APPLICATION NUMBER: JP 2001/246789
PRIOR FILING DATE: 201-66-18
NUMBER OF SEQ ID NOS: 2430
SOFTWARE: PATENTIN UNCL. 2.1
SEQ ID NO 636
LENGTH: 210
TYPE: PRT
      ; GENERAL INFORMATION:
                           ; Sequence 197, Application US/09886055; Patent No. US20020132273A1
                                                                    RESULT 3
US-09-886-055-197
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US-10-017-161-636
; Sequence 636, Application US/10; Publication No. US20030143668A1
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Best Local 9
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                                                                                                                                                              181
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                                                                                                                                                                                     PLIYSLRNKEVMRALRRVLGKYILLAHSTL 316
                                                                                                                                                                                                                                              RMPSNEGRKKALVTCSSHLIVVGMFYGAATFMYVLPSSFHSPKQDNIISVFYTIVTPALN
                                                                                                                                                                                                                                                                    RMPSNEGRKKALVTCSSHLIVVGMFYGAATFMYVLPSSFHSPKQDNIISVFYTIVTPALN 286
                                                                                                                                                          PLIYSLRNKEVMRALRRVLGKYILLAHSTL 210
                                                                                                                                                                                                                                                                                                                                  PFCVSWEIRHLLCEIPPLLKLACADTSRYELIIYVTGVTFLLLPISAIVASYTLVLFTVL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       66.5%; Score 210; DB 14; 100.0%; Pred. No. 3.3e-186; tive 0; Mismatches 0;
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               CURRENT APPLICATION NUMBER: US/09/804,291
CURRENT FILING DATE: 2001-03-13
PRIOR APPLICATION NUMBER: 60/188,914
PRIOR FILING DATE: 2000-03-13
PRIOR APPLICATION NUMBER: 60/192,033
PRIOR APPLICATION NUMBER: 60/192,033
PRIOR FILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: 60/198,474
PRIOR APPLICATION NUMBER: 60/199,335
PRIOR APPLICATION NUMBER: 60/199,335
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APPLICANT: Padigaru et al.

TITLE OF INVENTION: NOVEL:
PLUE REFERENCE: 11966-654

CURRENT APPLICATION NUMBER: US/09/777,789

CURRENT FILING DATE: 2001-05-11

PRIOR APPLICATION NUMBER: 60/180,646

PRIOR APPLICATION NUMBER: 60/180,646

PRIOR FILING DATE: 2000-02-07

NUMBER OF SEQ ID NOS: 101

SOFTMARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                              Sequence 197, Application US/09
Publication No. US20030088059A1
GENERAL INFORMATION:
APPLICANT: ZOZULA, SERGEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: STRYER, LUBERT
APPLICANT: ZOZULYA, SERGEY
TITLE OF INVENTION: RECEPTOR FINGERPRINTING, SENSORY PERCEPTION, AND
TITLE OF INVENTION: BIOSENSORS OF CHEMICAL SENSANTS
FILE REFERENCE: 078003-0277150
CURRENT APPLICATION NUMBER: US/09/806,055
CURRENT FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: 60/213,812
PRIOR APPLICATION NUMBER: 60/213,812
PRIOR PILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 522
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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Best Local Similarity
                                                                                                                                                                                                                     TITLE OF INVENTION: HUMAN OLFACTORY RECEPTORS AND GENES ENCODING SAME FILE REFERENCE: P 0278005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 316
TYPE: PRT
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ORGANISM: Homo sapiens
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  FILING
                                                                                                                                                                                                                                                                                                                                                                                                                                     52 EARLHMEMYLLIGQLSLMDLLFTSVVTPKALADFLRRENTISFGGCALQMFLALTMG 108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       h 18.0%; Score 57; DB 10; Length 316; Similarity 100.0%; Pred. No. 3.3e-44; 57; Conservative 0; Mismatches 0; Indels
DATE: 2000-04-24
                                                                                                                                                                                                                                                                                                                             Application US/09804291
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Gaps

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US-09-912-976-53
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                                                                                                                            ; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-912-976-53
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                                        Query Match
Best Local Similarity 100.
57; Conservative
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NUMBER OF SEQ ID NOS: 529
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 197
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Best Local S
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Smithson, Glennda
APPLICANT: Mishra, Vishnu
TITLE OF INVENTION: NOVEL PROTEINS AND NUCLEIC ACIDS ENCODING SAME
FILE REFERENCE: 21402-070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: APPLICANT:
                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/260,675
PRIOR FILING DATE: 2001-01-10
PRIOR APPLICATION NUMBER: 60/271,025
PRIOR FILING DATE: 2001-02-22
                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 2000-07-26
PRIOR APPLICATION NUMBER: 60/238,333
PRIOR FILING DATE: 2000-10-05
                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/912,976
CURRENT FILING DATE: 2001-07-05
PRIOR APPLICATION NUMBER: 60/221,336
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APPLICANT: Mezes, Peter
APPLICANT: Burgess, Cat
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PRIOR
                                                                                                                                                                                                SOFTWARE: P
                                                                                                                                                                                                                                  PRIOR FILING DATE: 2001-04-02
NUMBER OF SEQ ID NOS: 99
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                                                                                                                                                                               ENGTH: 316
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                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/278,164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 2000-09-07
APPLICATION NUMBER: 60/266,862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 2000-06-23
APPLICATION NUMBER: 60/226,534
FILING DATE: 2000-08-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 60/207,702
FILING DATE: 2000-05-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 60/230,732
FILING DATE: 2000-09-07
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52 EARLHMPMYLLLGQLSLMDLLFTSVVTPKALADFLRRENTISFGGCALQMFLALTMG
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                                                                                                                                                                                                                  PatentIn Ver. 2.1
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Gerlach, Valerie L
MacDougall, John R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alsobrook II, John P
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                                                                                                                                                                                                                                                                                     2001-03-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           William M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Muralidhara
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                                                     18.0%; Score 57;
100.0%; Pred. No.
tive 0; Mismatc
                                                                                                                                                                                                                                                                     60/280,876
                                                        Mismatches
                                             DB 11; __
3.3e-44;
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                                                                                       Length 316
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                                                      Indels
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                                                      Gaps
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PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: 60/237,581
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/238,735
PRIOR FILING DATE: 2000-10-06
PRIOR APPLICATION NUMBER: 60/240,736
PRIOR APPLICATION NUMBER: 60/240,736
PRIOR APPLICATION NUMBER: 60/260,019
PRIOR APPLICATION NUMBER: 60/260,019
PRIOR APPLICATION NUMBER: 60/260,338
PRIOR APPLICATION NUMBER: 60/260,338
                                                                                   Best Loc
Matches
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SEQ ID NO 58
LENGTH: 316
                                                                                                                              Query Match
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Publication No. US20030216545A1
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CURRENT APPLICATION NUMBER: US/09/965,422
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: 60/236,286
PRIOR FILING DATE: 2000-09-28
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APPLICANT: Tchernev, Velizar T
TITLE OF INVENTION: No. US20030:
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                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2001-02-26
NUMBER OF SEQ ID NOS: 127
                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2001-0: PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/262,156
PRIOR FILING DATE: 2001-01-17
PRIOR APPLICATION NUMBER: 60/262,498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Spytek, Kimberly A
                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/271,634
                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/263,133 PRIOR FILING DATE: 2001-01-19
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                                                                                                                                                                                      LENGTH: 316
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 2001-02-01
                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/263,691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2001-01-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2001-01-08
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                        52 EARLHMPMYLLLGQLSLMDLLFTSVVTPKALADFLRRENTISFGGCALQMFLALTMG 108
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                                                                                   Similarity
57; Conser
EARLHMPMYLLLGQLSLMDLLFTSVVTPKALADFLRRENTISFGGCALQMFLALTMG 108
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Ellerman, No...
Shlomit
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MacDougall, John
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Shenoy, Suresh G
Gerlach, Valerie
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Dickson, Kevin
Vernet, Corine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Smithson, Glennda
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                                                                                   18.0%; Score 57; DB 11;
larity 100.0%; Pred. No. 3.3e-44
Conservative 0; Mismatches (
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                                                                                                       3.3e-44;
                                                                                                                           Length 316;
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                                                                                   Gaps
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RESULT 8

US-10-343-650A-514

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Sequence 514, Application US/10343650A

Publication No. US20040067499A1

GENERAL INFORMATION:
APPLICANT: HAGA, TATSUYA

FILE REFERENCE: 31671-186347

CURRENT APPLICATION NUMBER: US/10/343,650A

CURRENT FILING DATE: 2003-07-21

PRIOR APPLICATION NUMBER: UF 2000/237818

PRIOR APPLICATION NUMBER: UF 2000/237818

PRIOR APPLICATION NUMBER: UF 2001/34434

PRIOR FILING DATE: 2001-02-13

NUMBER OF SEQ ID NOS: 694

PRIOR FILING DATE: 2001-02-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TYPE: PRT; ORGANISM: Homo sapiens US-10-343-650A-514
                                                                                  ; NAME/KEY: misc feature ; OTHER INFORMATION: Incyte ID No. US20030119111A1 7472446CD1 US-10-220-382-18
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SEQ ID NO 514
LENGTH: 316
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Publication No.
                                                                                                                                                                                                                               SOFTWARE: PERL Program SEQ ID NO 18
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                                         Query Match
                                                                                                                                                                                                                                                                  APPLICANT: BOROWSKY, Mark L.

TITLE OF INVENTION: GPROTEIN COUPLED RECEPTORS

FILE REFERENCE: PI-0044 PCT

CURRENT APPLICATION NUMBER: US/10/220,382

CURRENT FILING DATE: 2001-03-01

PRIOR APPLICATION NUMBER: 60/186,854; 60/188,384; 60/190,453; 60/190,730

PRIOR APPLICATION STEEL 2000-03-03; 2000-03-10; 2000-03-17; 2000-03-20

NUMBER OF SEQ ID NOS: 42
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APPLICANT:
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                                                                                                                                                                                        TYPE: PRT
                                                                                                                                                                ORGANISM: Homo sapiens
                                                                                                                                                                                                            ENGTH: 316
18.0%; Score 57; DB 14; Length 316; Local Similarity 100.0%; Pred. No. 3.3e-44; nes 57; Conservative 0; Mismatches 0; Indels
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LAL, Preeti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SHIH,
                                                                                                                                                                                                                                                                                                                                                                                                                   WALSH, Roderick T.
LO, Terence P.
BOROWSKY, Mark L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SHIH, Leo L.
TRIBOULEY, Catherine
LU, Dyung Aina M.
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YAO, Monique G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TANG, Y. Tom
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AU-YOUNG, Janice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             POLICKY, Jennifer L.
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hes 0;
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APPLICANT: SUWA, MAKIKO
APPLICANT: ASAI, KIYOSHI
APPLICANT: ASAI, KIYOSHI
APPLICANT: ASAI, KIYOSHI
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 084335/0152
CURRENT APPLICATION NUMBER: US/10/017,161
CURRENT FILING DATE: 2002-12-18
PRIOR APPLICATION NUMBER: JP 2001/246789
PRIOR FILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 2430
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                                                                                                             US-10-024-444B-3
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Query Match
Best Local Similarity
Matches 57; Conserv
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                                                                                                                                                                                                              SEQ ID NO 3
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Padigaru, Muralidhar
APPLICANT: Gerlach, Valerie L.
APPLICANT: Smithson, Glennda
                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Hart, Matthew
APPLICANT: Tomlinson, James E.
APPLICANT: Topper, James N.
APPLICANT: Kekuda, Ramesh
APPLICANT: Kekuda, Ramesh
APPLICANT: Kekuda, Ramesh
APPLICANT: Schlomit, Stacie J.
APPLICANT: Shlomit, Edinger R.
APPLICANT: Shlomit, Edinger R.
TITLE OF INVENTION: No. US20030165858Alel GPCR-Like Proteins
TITLE OF INVENTION: Same
FILE REFERENCE: 21402-224 AG
                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/256635
PRIOR FILING DATE: 2000-12-18
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/024,444B
                                                                                                                                    LENGTH: 316
TYPE: PRT
ORGANISM: human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stone, David
Bin-Yang, Ruey
Conley, Pamela B.
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          18.0%; Score 57; DB 14; Length 316; ilarity 100.0%; Pred. No. 3.3e-44; Conservative 0; Mismatches 0; Indels
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EARLHMPMYLLLGQLSLMDLLFTSVVTPKALADFLRRENTISFGGCALQMFLALTMG 108

52 52

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APPLICANT: MANG, KUN
APPLICANT: WANG, KUN
TITLE OF INVENTION: OLFACTORY RECEPTORS FOR ISOVALERIC ACID AND RELATED
TITLE OF INVENTION: MALODORANTS AND USE THEREOF IN ASSAYS FOR
TITLE OF INVENTION: MALODORANTS AND USE THEREOF IN ASSAYS FOR
TITLE OF INVENTION: IDENTIFICATION OF BLOCKERS OF MALODOR
TITLE OF INVENTION: UNMER: US/10/300,846
CURRENT APPLICATION NUMBER: 60/348,371
PRIOR APPLICATION NUMBER: 60/348,371
PRIOR APPLICATION NUMBER: 09/809,291
PRIOR FILING DATE: 2002-01-16
PRIOR APPLICATION NUMBER: 09/809,291
PRIOR APPLIC
                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/292,798
CURRENT FILING DATE: 2002-11-13
PRIOR APPLICATION NUMBER: 10/017,161
PRIOR FILING DATE: 2001-12-18
PRIOR APPLICATION NUMBER: JP 2001-246789
PRIOR PRIOR DATE: 2001-06-18
PRIOR FILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 2070
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 276
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US-10-300-846-26
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                                                                                                                                                                       ; TYPE: PRT; ORGANISM: Homo sapiens US-10-292-798-276
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APPLICANT: ASAI, KIYOSHI
APPLICANT: AKIYAMA, YUTAKA
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APPLICANT: ZOZULYA, SERGEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: ABURATANI, HIROYUKI
TITLE OF INVENTION: GUANOSINE TRIPHOSPHATE-BINDING PROTEIN COUPLED RECEPTORS
FILE REFERENCE: 084335/166
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            18.0%; Score 57; DB 15; Length 316; ilarity 100.0%; Pred. No. 3.3e-44; Conservative 0; Mismatches 0; Indels
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CURRENT APPLICATION NUMBER: US/09/974,591
CURRENT FILING DATE: 2001-10-09
PRIOR APPLICATION NUMBER: 60/245,292
PRIOR FILING DATE: 2000-11-02
NUMBER OF SEQ ID NOS: 26
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 14
LENGTH: 324
TYPE: PRT
ORGANISM: Homo sapiens
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SEQ ID NO 12
LENGTH: 324
TYPE: PRT
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Publication No. US20030059830A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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APPLICANT: Padigaru, Muralidhara
APPLICANT: Spytek, Kimberly A
TITLE OF INVENTION: No. US20030059830A1el Single Nucleotide Polymorphisms for Olfacto
TITLE OF INVENTION: Receptor-like Polypeptides and Nucleic Acids Encoding
TITLE OF INVENTION: the Same
FILLE REFERENCE: 15966-564 CIP
CURRENT APPLICATION NUMBER: US/09/974,591
CURRENT FILING DATE: 2001-10-09
PRIOR APPLICATION UNMERS: 60/245,292
PRIOR FILING DATE: 2000-11-02
NUMBER OF SEG ID NOS: 26
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APPLICANT: Burgess, Catherine E
APPLICANT: Grosse, William M
                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Alsobrook II, John P
APPLICANT: Burgess, Catherine E
APPLICANT: Grosse, William M
APPLICANT: Lepley, Denise M
APPLICANT: Padigaru, Muralidhara
APPLICANT: Spytek, Kimberly A
APPLICANT: Syytek, Kimberly A
APPLICANT: NOENTION: No. US20030059830Alel Single Nucleotide Polymorphisms for Olfacts
TITLE OF INVENTION: Receptor-like Polypeptides and Nucleic Acids Encoding
TITLE OF INVENTION: the Same
FILE REFERENCE: 15966-654 CIP
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tive 0; Mismatches 0;
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score greater than or equal to the score of the result being printed,
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AG090554 Pan trogl
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00 CH230-47	מוזיסובה השישור	72978 OR 2	171253 OR 20	B173653 OR 20	3224 OR_2c	B173469 OK_	656752 AGE	7/338 Pan	2220 22	CORL	8 / /3 L CD /	74/21 CIVATO 24/4/	/4185 BY6	9244 AL53	931094 RPC1-4	/41/8 IMULZE	1 NO.	73379 OR 202	3,5	202	2-2-10	E6446/1 Cigr-988	KU16560 Mus musc	17036 Mus	BY716264 BY716264	92351	42411	73558	W431124	7342	I399100 MI-I	49576 CH24	191 71360 M	85 262681	Z709687 RPCI-2	352344 CH230	Z323827 1M0045	B173529 OR 202	B174035 OR_2	Z806221 2M0068E1	Z373068 1M012

ALIGNMENTS

031161 CUS	AG031161 710 bp DNA linear GSS 01-NOV-2001
FINITION CESSION	Pan troglodytes DNA, Cione: Fis-Vosavi.E, Schoule Burie, Cognomic AG031161
RSION	AG031161.1 GI:16558034
YWORDS	GSS.
URCE	Pan troglodytes (chimpanzee)
ORGANISM	Pan troglodytes
	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buterescomi;
	Mammalia; Eutheria; Primates; Catarrnini; Hominicae; Faii.
FERENCE	
AUTHORS	Fujiyama, A., Hattori, M., Toyoda, A., Tayror, 1.D., rada, 1.,
	Totoki,Y., Watanabe,H. and Sakaki,I.
TITLE	BAC end sequences of Library PTB
JOURNAL	Unpublished
FERENCE	2 (bases 1 to 710)
AUTHORS	Fujiyama,A., Hattori,M., Toyoda,A., Tayror,Toroki V., Watanabe,H. and Sakaki,Y.
TITLE	Direct Submission
JOURNAL	Submitted (02-AUG-2001) Asao Fujiyama, The institute of rhysical
	and Chemical Kesearch (Kindw), Genomic Beterros 200-20-1-7-7-7-7-7-7-7-7-7-7-7-7-7-7-7-7-7-7-
	(E-mail:chimpbes@gsc.riken.go.jp, URL:http://hgp.gsc.riken.go.jp/,
	Tel:81-45-503-9111, Fax:81-45-503-9170)
OMMENT	Clones are derived from the Chimpanzee BAC Library Fib III Division of
	Mas Actigiacen mutify and the Francisco

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                                                                                                                                                                                                                                                                                                                                                                            AUTHORS
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                                                                                                   Submitted (02-AUG-2001) Asao Fujiyama, The Institute of Physical and Chemical Research (RIKEN), Genomic Sciences Center (GSC); 1-7-22 Suehizo-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan (E-mail:chimpbes@gsc.riken.go.jp, URL:http://hgp.gsc.riken.go.jp/, Tel:81-45-503-9111, Fax:81-45-503-9170)

Clones are derived from the chimpanzee BAC library PTB This BAC end was generated during the R&D process and may have higher chance of nptwers.
                                                                                                                                                                                                                                                                                                                                Fujiyama,A., Hattori,M., Toyoda,A., Totoki,Y., Watanabe,H. and Sakaki,Y. BAC end sequences of Library PTB
                                                                                                                                                                                                                                                            Fujiyama, A., Hattori, M., Toyoda, A., Totoki, Y., Watanabe, H. and Sakaki, Y.
                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
                                                                                                                                                                                                                                               Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                           Pan troglodytes (chimpanzee)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AG090554
AG090554.1 GI:16642356
GSS.
                                                       Sequencing: M13Rev
                                                                                                                                                                                                                                                                                                                       Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pan troglodytes DNA, clone:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequencing:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGCTTATAATATACGTGACAGGTGTGACTTTCCTCTTGCTCCC 648
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGAGCCCAAGAGTCTGCTGGATCATGGTGGCCACATCCTGGATCCTGGCATCCCTGATTG 485
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TGAGCCCAAGAGTCTGCTGGATCATGGTGGCCACATCCTGGATCCTGGCATCCCTGATTG 538
                                                                                                                                                                                                                                                                                                    (bases 1 to 663)
 R.Site 1
R.Site 2
                                    Vector
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="genomic DN;
/db_xref="taxon:9598"
/clone="PTB-005A07.F"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /cell_type="lymphoblast"
/clone_lib="PTB Chimpanzee Male BAC Library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Pan troglodytes"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sex="male"
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: pKS145
: SacI
: SacI.
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: SacI
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Pred. No. 3.2e-52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    663
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 bp DNA
PTB-090C09.R,
                                                                                                                                                                                                                                                                            Taylor, T.D., Yada, T.,
                                                                                                                                                                                                                                                                                                                                                                     Taylor, T.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear GSS 03-NOV-200 genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 710;
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FEATURES
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AUTHORS
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SOURCE
ORGANISM
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AQ522745
LOCUS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JOURNAL
MEDLINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PUBMED
                                                                                                                                                                                              source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            233 CTGTTCACATCTGTTGTCACTCCCAAGGC 261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            401 Queen Anne Avenue North, Seattle, WA 98109, USA
Tel: (206) 616-3618
Fax: (206) 616-368
Fax: (206) 616-3687
Email: jwallace@u.washington.edu
Clones are derived from the human BAC library RPCI-11. For BAC
library availability, please contact Pieter de Jong
(pieter@dejong.med.buffalo.edu). Clones may be purchased from
BACPAC Resources (http://bacpac.med.buffalo.edu/ordering bac.htm)
or from Resear h Genetics (info@resgen.com). BAC end Web Server:
                                                                                                                                                                                                        High quality sequence stop: 566.
Location/Qualifiers
                                                                                                                                                                                                                                                                          Seq primer: T7
                                                                                                                                                                                                                                                                                           http://www.htsc.washington.edu
Plate: 797 row: D column: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High Throughput Sequencing Center University of Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 566)

Mahairas,G.G., Wallace,J.C., Smith,K., Swartzell,S., Holzman,T., Keller,A., Shaker,R., Furlong,J., Young,J., Zhao,S., Adams,M.D. at uncall.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AQ522745

AQ52271B1 B10 T7A RPCI-11 Human Male BAC Library Homo sapiens genomic clone Plate=797 Col=19 Row=D, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Mahairas GG, Wallace JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             scanning the human genome Proc. Natl. Acad. Sci. U.S.A. 96 (17), 9739-9744 (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence-tagged connectors: A sequence approach to mapping and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .68
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Conservative (
/clone lib="RPCI-11 Human Male BAC Library" /note="Vector: pBACe3.6; Site 1: EcoRI; Site 2: EcoRI; Male blood DNA was isolated from one randomly chosen do and partially digested with a combination of EcoRI and
                                                                                                                            /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                       , . . 566
                                                                                        sex="male"
                                                                                                           clone="Plate=797 Col=19 Row=D"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Pan troglodytes"
/mol_type="genomic DNA"
/db_xref="taxon:9598"
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/clone_lib="PTB Chimpanzee Male BAC Library"
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100.0%; Pred. No.
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RESULT 4
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ORGANISM
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VERSION
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                                RESULT 5
AZ373068/c
DEFINITION
                  Locus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AUTHORS
TITLE
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Best Local Similarity
                                                                                                                                                                    Best Local Similarity
Matches 43: Common
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JOURNAL
                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                       source
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                                                                                                                                         634
                                                                                                       129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BE897401 843 bp mRNA linear EST 20-OCT-2000 BE897401 NIH_MGC_72 Homo sapiens cDNA clone IMAGE:3922342 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM9756 row: 1 column: 23
High quality sequence start: 12
High quality sequence stop: 614.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 843)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mRNA sequence.
 AZ373068 625 bp DNA linear GSS 02-OCT-20-
1M0125N08F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BE897401.1 GI:10362826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BE89740
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium DNA Sequencing by: Incyte Genomics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTACATTTACAATCCTATACATGT 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTACATTTACAATCCTATACATGT 515
                                                                                                                       TITCCTCTTGCTCCCATTTCTGCCATTGTGGCCTCCTACACA 676
                                                                                                                                                           4.4%; Score 43; ilarity 100.0%; Pred. No. Conservative 0; Min-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EcoRI Methylase. Size selected DNA was cloned into the pBACe3.6 vector at EcoRI sites"
                                                                                                                                                                                                                                                           /tissue type="melanotic melanoma"
/lab_host="DH10B (phage-resistant)"
/clone lib="NIH MGC 72"
/note="Organ: skin; Vector: pcMV-SPORT6; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 2 kb. Library constructed by Life Technologies."
                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Homo sapiens"

/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                    /db_xref="taxon:9606"
/clone="IMAGE:3922342"
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; Pred. No. 1.2e-32
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                                                                                                                                                                                            6.3e-11;
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                                                                                                                                                                                                            Length 843;
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2M0068E10F Mouse 10kb plasmid
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bp DNA linear GSS 2

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Query Match
Best Local Similarity
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Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome C.
University of Utah
255 ATGGCCTATGACAGGTATGTGGCCATTTGTCATCCTCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb
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Mammalia; Eutheria; Rodentia;
1 (bases 1 to 625)
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High quality sequence stop: 625.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0125 row: N column: 08
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                      38;
                                                ATGGCCTATGACAGGTATGTGGCCATTTGTCATCCTCT
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                                                                                                                                                                                                                  was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/clone="UUGC1M0125N08"
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strain="C57BL/6J"
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                                                                                                                             100.0%;
                                                                                                                             3.9%; Score 38;
100.0%; Pred. No.
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Sciurognathi; Muridae; Murinae; Mus
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  CB174035 528 bp mRNA
OR_2032D01_010529.yl Adult mouse olfactory
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Insert Length: 10000 Std Error: 0.00
Plate: 0068 row: E column: 10
Seq primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plasmid inserts
Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 801 585 5606
Fax: 801 585 7177
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AZ806221
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                                                                                                                                                                                                                                                                                                                                                                 adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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strain="C57BL/6J"
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Pred. No. 0.0001;
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                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 595)
Young,J.M., Shykind,B.M., Lane,R.P., Tonnes-Priddy,L., Ross,J.A., Walker,M., Williams,E.M. and Trask,B.J.
Odorant receptor expressed sequence tags demonstrate olfactory expression of over 400 genes, extensive alternate splicing and
                                                                                                                                                                                                                                                                                                                                               CB173529 595 bp mRNA linear EST 09-OCT-200 OR 2028D09 010501.yl Adult mouse olfactory epithelium library Musmusculus cDNA clone 2028D09 5', mRNA sequence.
                 unequal expression levels
Genome Biol. 4 (11), R71.1-R71.15
                                                                                                                                                                                                                                      Mus musculus
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CB173529.1 GI:37592158
                                                                                                                                                                                                                                                             Mus musculus (house mouse)
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Genome Biol. 4 (11), R71.1-R71.15 (2003)
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Seq primer: M13 Reverse.
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Young gene new name GA_x6K02T2Q125-47640742-47639798
Young gene old name GA_x5J8B7W3KVV-147900-148844
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Fred Hutchinson Cancer Research Center
1100 Fairview Avenue N., C3-168, P.O. Box 19024, Seattle, WA
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Fax: 206 667 6524
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Young JM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 528)
Young,J.M., Shykind,B.M., Lane,R.P., Tonnes-Priddy,L., Ross,J.A., Walker,M., Williams,E.M. and Trask,B.J.
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Mus musculus
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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/dev_stage="Adult"
/clone_lib="Adult"
/clone_lib="Adult mouse olfactory epithelium library"
/note="Torgan: Olfactory turbinates; Vector:
LambdaZAPII-XR; Site_1: EcoRI; Site_2: XhoI; This library
was provided by Leslie Vosshall. mRNA was prepared from
the olfactory and respiratory epithelium of an adult
mouse. Oligo-dT primed cDNA was directionally cloned into
Stratagene's lambdaZAPII-XR vector."
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/mol_type="mRNA"
/db_xref="taxon:10090"
/clone="2032D01"
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AUTHORS
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Fred Hutchinson Cancer Research Center
1100 Fairview Avenue N., C3-168, P.O. F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zhang gene name MOR185-4
Seq primer: M13 Reverse.
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Young gene new name GA_x6K02T2Q125-47640742-47639798
Young gene old name GA_x5J8B7W3KVV-147900-148844
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 206 667 1471 Fax: 206 667 6524
                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Insert O045 row: B column: 03
Seg primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 432)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
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432 bp DNA linear GSS 29-SEP-201
1M0045B03R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
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                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                         University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                       Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                       Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                             plasmid
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                                                                         High quality sequence stop: 432.
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/clone_lib="Adult mouse olfactory epithelium library"
/note="Organ: Olfactory turbinates; Vector:
LambdaZAPII-XR; Site_1: EcoRI; Site_2: XhoI; This library
was provided by Leslie Vosshall. mRNA was prepared from
the olfactory and respiratory epithelium of an adult
mouse. Oligo-dT primed cDNA was directionally cloned into
Stratagene's lambdaZAPII-XR vector."
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/organism="Mus musculus"
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/mol_type="mRNA"
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Pred. No. 0.00013;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Shaying Zhao
Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville,
Tel: 301 838 0200
Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rattus norvegicus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BH352344 545 bp DNA linear (CH230-136E3,TV CHORI-230 Segment 1 Rattus norvegicus
                                                                                  Clones are derived from the rat BAC library CHORI-230 (http://www.chori.org/bacpac/rat230.htm). For BAC library availability, please contact Pieter de Jong (pdejong@mail.cho.org). Clones may be purchased from BACPAC Resources (http://www.chori.org/bacpac/or ering_information.htm). BAC end page: http://www.cigr.org/tdb/bac_ends/rat/bac_end_intro.html plate: 136 row: E column: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zhao,S., Shetty,J., Shatsman,S., Tsegaye,G., Geer,K., Shvartsbeyn,A., Gebregeorgis,E., Overton,L., Russell,D., Chen,D., Riggs,F., de Jong,P. and Fraser,C.M.
Rat BAC End Sequences from Library CHORI-230 EcoRI segment
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Seq primer: T7
Class: BAC ends.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rattus norvegicus (Norway rat)
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Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adaptored DNA was purified and size-selected for a 9.5 to
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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Locus

KEYWORDS VERSION

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FEATURES

Location/Qualifiers

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Best

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AUTHORS
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AZ709687/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      369 TCATGGCCTATGACAGGTATGTGGCCAT 396
                                                                                                                                                                                                                                                                                                                                                                                                                      Email: szhao@tigr.org
Clones are derived from the mouse BAC library RPCI-24. For BAC
library availability, please contact Pieter de Jong
(pdejong@mail.cho.org). Clones may be purchased from BACPAC
Resources (http://www.chori.org/bacpac/orderingframe.htm). BAC (
page: http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html
Plate: 82 row: P column: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zhao, S., Nierman, W., Malek, J., Shatsman, S., Akinret, B., Levins Tsegaye, G., Geer, K., Krol, M., Shvartsbeyn, A., Gebregeorgis, E., Russell, D., de Jong, P. and Fraser, C.M.
Mouse BAC End Sequences from Library RPCI-24
Unpublished (1999)
Other GSSs: RPCI-24-82P2.TV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28;
                                                                                                                                                                                                                                                                                                                                                                                          Class: BAC ends.
                                                                                                                                                                                                                                                                                                                                                                                                                 Seq primer: SP6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 301 838 0200
Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The Institute for Genomic Research
9712 Medical Center Dr., Rockville, MD 20850, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Department of Eukaryotic Genomics
The Institute for Genomic Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Shaying Zhao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mammalia; Eutheria;
1 (bases 1 to 680)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AZ709687 680 bp DNA linear GSS 24-JAN-2001 RPCI-24-82P2, TJ RPCI-24 Mus musculus genomic clone RPCI-24-82P2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.9%; Score 28; llarity 100.0%; Pred. No. Conservative 0; Mismatcl
                      /cell type="Spleen/Brain"
/celne lib="RPCI-24"
/clone lib="RPCI-24"
/clone lib="RPCI-24"
/note="Vector: pTARBAC1; Site 1: BamH1; Site 2: BamH1;
/note="Vector: pTARBAC1; Site 1: BamH1; Site 2: BamH1;
RPCI-24 Mouse BAC Library produced by Pieter de Jong. The
library was cloned in the pTARBAC1 cloning vector at the
library was cloned in the pTARBAC1 cloning vector at the
BamH1 sites using MboI partially digested male C57BL/6J
                                                                                                                                                                                                                                                              /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /cell_type="Brain"
/clone lib="CHORI-230 Segment 1"
/clone="Vector: pTARBAC2.1; Site_1: EcoRI; Site_2: EcoRI;
CHORI-230 Rat (BM/SsNHsd/MCW) BAC library produced by
Pieter de Jong"
                                                                                                                                                                                                                /db_xref="taxon:10090"
/clone="RPCI-24-82P2"
                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                            /sex="Male"
                                                                                                                                                                                                                                                                                                              organism="Mus musculus"
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/strain="BN/SsNHsd/MCW"
/db_xref="taxon:10116"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="CH230-136E3"
/sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Rattus norvegicus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 28; Length 545; 0.005;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Levins, M.,
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RESULT 13
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ORGANISM
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ACCESSION
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BF444685
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      KEYWORDS
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Matches
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MEDLINE
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Best Local !
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                                                                                                                                                                                                                                                                                                                                          881 CTCATCTACAGCCTGAGGAATAAGGA 906
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AW486191 334 bp mRNA
71360 MARC 1PIG Sus scrofa cDNA 5', mRNA
AW486191 GI:7056297
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BACKWARD: GTTTTCCCAGTCACGACG Plate: 98 row: G column: 8 seq primer: ATTTAGGTGACACTATAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Smith TPL USDA, ARS, US Meat Animal Research Center DSDA, ARS, US Meat Animal Research Center PO Box 166, Clay Center, NE 68933-0166, USA Tel: 402 762 4366

Fax: 402 762 4390

Email: smith@email.marc.usda.gov

Single pass sequencing. Bases called and alt trimmed with phred v0.980904.e. Vector identified by cross_match with the -minscore and -minmatch 12 options.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 222)
Fahrenkrug, S.C., Smith, T.P.L., Freking, B.A., Cho, J., White, J., Vallet, J., Wise, T., Rohrer, G.A., Pertea, G., Sultana, R., Quackenbush, J. and Keele, J.W.
Porcine gene discovery by normalized cDNA-library sequencing and EST cluster assembly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EST.
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Mammalia; E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sus scrofa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sus scrofa
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BF444685.1 GI:11504777
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BF444685
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CATGGCCTATGACAGGTATGTGGCCATT 397
                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genome 13 (8), 475-478 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="unvn | /mol_type="unvn | /mol_type="pooled" | /tissue_type="pooled" | /lab_host="DH10B" | /clone_lib="marc_2pig" | /clone_lib="marc_2pi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Vector: pCMV SPORT6; Site 1: NotI; Site 2: SalI; Library made from pooled tissue from testis, ovary, endometrium, hypothalamus, pituitary, and placenta."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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100.0%; Pr
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100.0%; Pred. No.
tive 0; Mismatc
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 26; DB 10;
Pred. No. 0.046;
                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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RESULT 14
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Best Local
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                                                                                         JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         90
Unpublished Contact: Harris Lewin
Contact: Harris Lewin
Department of Animal Sciences
University of Illinois at Urbana Champaign
University of Transcript Urbana, IL 61801, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26;
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1 (bases 1 to 334)

Pahrenkrug,S.C., Smith,T.P.L., Freking,B.A., Cho,J., White,J., Vallet,J., Wise,T., Rohrer,G.A., Pertea,G., Sultana,R., Quackenbush,J. and Keele,J.W.

Porcine gene discovery by normalized cDNA-library sequencing and Porcins assembly
                                                                                      Genome Sequence
Unpublished (2003)
                                                                                                                                          Larkin,D.M., Everts-van der Wind,A., Rebeiz,M., Bachman,S., Green,S., Campos,E.J., Benson,L.D., Womack,J.E., de Jong,P.J. and Lewin,H.A.
                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                      genomic survey
BZ949576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and trimmed with phred
v0.980904.e. Vector identified by cross_match with the -mins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Smith TPL USDA, ARS, US Meat Animal Research Center PO Box 166, Clay Center, NE 68933-0166, USA Tel: 402 762 4366 Fax: 402 762 4390
                                                                                                                                                                                                                                                                           Bos taurus
Bos taurus
                                                                                                                                                                                                                                                                                                                 GSS
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                                                                                                                           A Cattle-Human Comparative Map Built with Cattle BAC-ends and Human
                                                                                                                                                                                                                                                                                                                                  BZ949576.1 GI:31734881
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Seq primer: ATTTAGGTGACACTATAG.
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/note="Vector: pCMV SPORT6; Site 1: NotI; Site 2: SalI;
/ibrary made from pooled tissue from day 11, 13, 15, 20,
and 30 embryos."
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/lab_host="DH10B"
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0.051;
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                                                                                                                                                                                  Schweitzer, P.,
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                                                                                                                                                                Liu, L.,
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KEYWORDS
SOURCE
ORGANISM
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BI399100/c
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MEDLINE
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                                                                                                                                                                                                                                                                                                                                                                                                                               AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: h-lewin@uiuc.edu
Clones are derived from the bovine BAC library CHORI-240
(http://www.chori.org/bacpac/bovine240.htm). For BAC library
availability, please contact Pieter de Jong (pdejong@mail.cho.org).
Clones may be purchased from BACPAC Resources
(http://www.chori.org/bacpac/ordering_information.htm). This work
was undertaken as part of the International Bovine BAC Mapping
Consortium (IBBMC) by by University of Illinois at Urbana
Champaign, USA with funds provided by grant No. AG202-34480-11828
from USDA-CSREES and AG99-35205-8534 from USDA/NRI (Livestock
Genome Sequencing Initiative)
Plate: 38 row: O column: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BI399100 451 bp mRNA linear EST MI-P-AY1-nrk-c-11-0-UI.s1 MI-P-AY1 Sus scrofa cDNA clone MI-P-AY1-nrk-c-11-0-UI 3', mRNA sequence.
oligonucleotide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail. The sequence tag present in the cDNA between the NotI site and the oligo-dT track served to verify it as a clone from the normalized placenta library cDNA Library Preparation: M.B. Soares Lab, University of Iowa EST sequencing: M.B. Soares Lab, University
                                                                                                                                                                                                 Contact: Tuggle CK
Molecular Genetics Laboratory, Department
Iowa State University
201 Kildee Hall, Ames, IA 50011-3150, USA
Tel: 5152944252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sus scrofa (pig)
Sus scrofa
                                                                                                                                                                                                                                                                                                                      8889548
                                                                                                                                                                                                                                                                                                                                                             Genome Res. 6 (9),
                                                                                                                                                                                                                                                                                                                                                                                       discovery
                                                                                                                                                                                                                                                                                                                                                                                                                            Bonaldo, M.F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae;
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Class: BAC ends.
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                                                                                                                                    The sequence contained an oligo-dT track that was present in the
                                                                                                                                                       Email: cktuggle@iastate.edu
                                                                                                                                                                                   Fax: 5152942401
                                                                                                                                                                                                                                                                                                                                                                                                          Normalization and subtraction: two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    (bases 1 to 451)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="Vector: pTARBAC1.3; Site 1: MboI; Site 2: MboI; Hereford bull L1 Domino 99375; CHORI-240 Bovine BAC library (Male) produced by Pieter de Jong"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone="CH240_3809"
/sex="Male"
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/strain="breed: Hereford"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GI:15178161
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               _type="Blood"
=_lib="CHORI-240"
                                                                                                                                                                                                                                                                                                                                                                791-806 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 26;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                               and Soares, M.B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                       Animal
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Sus.
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Seq primer: M13 Foorward
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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (03-SEP-2002) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA On Sep 3, 2002 this sequence version replaced gi:22123080.
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Smit, A.F.A. & Green, P. (1996-1997)
http://ftp.genome.washington.edu/RM/RepeatMasker.html
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------ Project Information
Center project name: Li1860
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (11-JUL-2001) Makiko Suwa, Computational Biology Research Center (CBRC), National Institute of Advanced Industrial Science and Technology (AIST); 2-41-6 Aomi Koto-ku, Tokyo 135-0064, Japan (E-mail:m-suwa@aist.go.jp, URL:http://www.cbrc.jp/, Tel:B1-3-3599-8080, Fax:B1-3-359-8081) Tel:B1-3-3599-8080, Fax:B1-3-359-8081) Tel:B1-3-359-8080, Fax:B1-3-359-8081, Fax:B1-
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                                                                                                                             TGGAGGCTGTGCACTTCAGATGTTCCTGGCACTGACAATGGGTAGCGCTGAGGACCTCCT 360
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llarity 99.9%;
Conservative
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/mol_type="genomic DNA"
/isolate="CBRC7TM_102"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /product="seven transmembrane helix receptor"
/evidence=not_experimental
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AUTHORS
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AC027641/c
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JOURNAL
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           421
                                                                                                                                                                                   Submitted (30-MAR-2000) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St. MO 63108, USA
On Jun 17, 2000 this sequence version replaced gi:7344780.
                                                                                                                                                                                                                                                                                      Direct Submission
                                                                                                                                                                                                                                                                                                                2 (bases 1 to 208430) Waterston, R.H.
                                                                                                                                                                                                                                                                                                                                                           Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
1 (Dases 1 to 208430)
Waterston, R.H.
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ACO27641.2 GI:8570385
HTG: HTGS_PHASE1; HTGS_DRAFT.
Homo sapiens (human)
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ACO27641

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Sequencing vector: plasmid; 2%
Sequencing vector: plasmid; 2%
Chemistry: Dye-primer ET; 98% of reads
Chemistry: Dye-terminator Big Dye; 2% of reads
Assembly program: Phrap; version 0.990319
Consensus quality: 19825 bases at least Q40
Consensus quality: 202103 bases at least Q30
Consensus quality: 203934 bases at least Q20
Insert size: 219000; agarose-fp
Insert size: 206730; sum-of-contigs
Quality coverage: 4.27 in Q20 bases; agarose-fp
Quality coverage: 4.55 in Q20 bases; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NOTE: This is a 'working draft' sequence. It currently consists of 18 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence.
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                                                                    /note="assembly_name:Contig10"
5729. .7760
                                                                                                               1861. .3171
/note="assembly_name:Contig9"
3272. .5628
'note="assembly_name:Contig12'
                          /note="assembly_name:Contig11"
7861. .15503
                                                                                                                                                                            /note="assembly_name:Contig8"
                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
/chromosome="11"
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                                                                                                                                                                                                                           /clone="RP11-732A19"
                                                                                                                                                                                                                                                                                                                                                                            87730: gap of unknown length
106802: contig of 19072 bp in 10
106902: gap of unknown length
120650: contig of 13748 bp in 10
120750: gap of unknown length
156637: contig of 35887 bp in 10
156737: gap of unknown length
208430: contig of 51693 bp in 10
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1860: gap of unknown length
3171: contig of 1311 bp in length
3271: gap of unknown length
5628: contig of 2357 bp in length
5728: gap of unknown length
7760: contig of 2032 bp in length
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                          GATTGCTATAGGACATACCATGTACACTATGCACCTCCCTTTCTGTGTGTCCTGGGAAAT
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CAGGCATCTGCTCTGTGAGATCCCACCCTTGCTGAAGTTGGCCTGTGCTGATACCTCCAG
                                          CCTCATGAGCCCAAGAGTCTGCTGGATCATGGTGGCCACATCCTGGATCCTGGCATCCCT
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120751. .156637
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73854. .87630
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Patent: WO 0127158-A 430 19-APR-2001;
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AX241682
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        CTCCTGCTTGGGCAGCTCTCTCATGGACCTCCTGTTCACATCTGTTGTCACTCCCAAG
                                                     AGCAATGGTCTGCTGCTCCTGGCCATCACCATAGAAGCCCGGCTCCACATGCCCATGTAC
                                     AGCAATGGTCTGCTGCTGGCCATCACCATAGAAGCCCGGCTCCACATGCCCATGTAC
                                                                                                                                                                                                          Conservative
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/mol type="unassigned DNA"
/mb xref="teaxon:32630"
/noTe="(H38g279 nucleotide)"
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                                             Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi, Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 649)

Fuchs, T., Malecova, B., Linhart, C., Sharan, R., Khen, M., Herwig, R., Shmulevich, D., Elkon, R., Steinfath, M., O'Brien, J.K., Radelof, U., Ehrach, H., Olender, Z., Glusman, G., Lancet, D. and Shamir, R. DEFOG: A Practical Scheme for Deciphering Families of Genes
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       Linhart, C., ., Steinfath,
     Sharan,R., Khen,M., Herwig,R.,M., O'Brien,J.K., Radelof,U.,
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                                                                                                               Euteleostomi; Homo.
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Direct Submission
Submitted (16-JUL-2001) Dept. of Molecular
Human Genome Center, Weizmann Institute of
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/note="contains transmembr
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Patent: WO 0127158-A 427 19-APR-2001;
Digiscents (US) ; YEDA RESEARCH AND DEVELOPMENT
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                         Lal,P., Tang,Y.T., Patterson,C., Yao,M.G., Shih,L.L., Tribouley,C.M., Lu,D.A., Yue,H., Khan,F.A., Policky,J.L., Au-Young,J., Yang,J., Harland,L., Walsh,R.T., Lo,T.P. and Borowsky,M.L.
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Patent: WO 0166742-A 39 13-SEP-2001;
Incyte Genomics, Inc. (63)
 AX448455
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/mol type="unassigned DNA"
/mol type="taxon:32630"
/mbe="(H38g276 nucleotide)"
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                                                                                                                                                                                                                                                                        /note="Incyte ID No: 7472446CB1"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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1 (bases 1 to 951)

Haga, T., Takeda, S. and Miyake, N.
Novel G-protein coupled receptors
Patent: JP 2002112793-A 257 16-APR-2002;
JAPAN SCIENCE AND TECHNOLOGY CORP
OS Homo sapiens (human)
PN JP 2002112793-A/257
                                                                                                                                                                                                                                                                                                                                                                                      BD144532.1 GI:27850290 JP 2002112793-A/257.
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G01N33/566//
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C C12N5/10,C12Q1/02,C12Q1/68,G01N33/15,G01N33/50,G01N33/53, PC
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/ CAD XYEE = "REINERMEL CAD 37547"
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/ CAD XYEE = "REINIEMBL CAD 37547"
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      Eukaryota; Metazoa;
Mammalia; Eutheria;
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AX551018
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Baumgartner, J.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Novel single nucleotide polymorphisms for olfactory receptor-like
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AX210248
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 275 from Patent EP1270724.
AX646083
AX646083.1 GI:28798462
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                                                    Suwa,M., Asai,K., Akiyama,Y. and Aburatani,H.
Guanosine triphosphate-binding protein coupled receptors
Patent: EP 1270724-A 275 02-JAN-2003;
National Institute of Advanced Industrial Science and Technology
(JP); Center for Advanced Science and Technology Incubation, Ltd.
                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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larity 100.0%; Pred. No. 2.8e-58;
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/mol_type="unassigned DNA"
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/protein_id="R2A969177.1"
/protein_id="R2A969177.1"
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/LLALTMERALHMPMYLLGG(ISLMDILFTSVUTPKALADFLRENTISFGGCALQMF
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/YTMIYPFCBAQEIRHLLCEIPHLLKVACADTSRYELMYVMGVTFLLPSLAAILASY
/TQILLTYLHMPSNEGRKKALVTCSSEHLTVVCMPYGGAATFMYVLPSSFHSTRQDNIISV
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ALIGNMENTS

RESULT 1
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AC ABQ81238 GPCR1; olfactory receptor; G-protein coupled receptor; receptor; human; antiarteriosclerotic; antidiabetic; analgesic; cytostatic; antidepressant; nootropic; neuroprotective; immunosuppressive; antiallergic; antibacterial; virucide; protozoacide; fungicide; cardiomyopathy; atherosclerosis; diabetes; gene therapy; gene; ss. Homo sapiens. WO200270707-A2 Human olfactory receptor-like protein GPCR1 coding sequence. 05-DEC-2002 ABQ81238 standard; cDNA; 977 BP. (first entry) Location/Qualifiers 20. .970 /*tag= a /product= "GPCR1"

18-DEC-2000; 2000US-0256635P. 18-DEC-2001; 2001WO-US049168. CURAGEN CORP.
COR THERAPEUTICS INC.

Padigaru M, Gerla Conley P, Hart M, Edinger S, Macdou Macdougall JR; Gerlach VL, Smithson G, Stone D, Bin-Yang art M, Tomlinson JE, Topper JN, Kekuda R, R; Casman SJ;

WPI; 2002-698753/75. P-PSDB; ABB79908.

New G-protein coupled receptor (GPCR1) proteins and nucleic acids, useful for treating or preventing a GPCR1-associated disorder, e.g. cardiomyopathy, atherosclerosis, diabetes, or disorders related to cell signal processing.

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The present sequence is that of a nucleic acid encoding a novel human G-CC protein coupled receptor, designated GPCR1, that has structural and CC physiological functions characteristic of the olfactory receptor family. The invention provides GPCR1 polypeptides, nucleic acids and antibodies. The invention provides GPCR1 polypeptides, nucleic acids and antibodies. The invention provides GPCR1 polypeptides, nucleic acids and antibodies. CC claimed in processing or metabolic pathway modulation in humans cC (claimed). They may further be used for treating or preventing e.g. CC (claimed). They may further be used for treating or preventing e.g. CC (claimed). They may further be used for treating or preventing e.g. CC (claimed). They may further be used for treating or preventing e.g. CC (claimed). They may further be used for treating or preventing e.g. CC (claimed). They may further be used from the protein and inventy or schizophrenia), cc psychotic and neurological disorders (e.g. anxiety or schizophrenia), and CC anteriors, neurodegenerative diseases (e.g. Alzheimer's disease), and CC infections, neurodegenerative diseases (e.g. Alzheimer's disease), and CC as research tools, as diagnostic or prognostic markers, in gene therapy, as research tools, as diagnostic or prognostic markers, in gene therapy, as research tools, as diagnostic or prognostic markers, and as biological collections weapons. The nucleic acids may also be used in chromosome conditions of the surface of a biological sample, as surrogate markers, and in pharmacogenomics of a biological sample, as surrogate markers, and in pharmacogenomics of the prognostic markers and in pharmacogenomics prognostic markers and in pharmacogenomics of the prognostic marke
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Claim 8; Page 352; 1857pp;

English.

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CAGGCATCTGCTCTGTGAGATCCCACCCTTGCTGAAGTTGGCCTGTGCTGATACCTCCAG

GTATGAGCTTATAATATACGTGACAGGTGTGACTTTCCTCTTGCTCCCCATTTCTGCCAT

660 600 600 540

The present sequence is one of a number of isolated polynucleotides which encode polypeptides involved in olfactory sensation. The polynucleotides can be used in screening for olfactory agonists and antagonists. The methods allow for the determination of primary scents and the scents. The methods also enable determination of secondary scents and the scents. The methods also enable determination of secondary scents and the identification of combinations of odour receptors that are involved in detecting such secondary scents. This enables the construction of a scent representation (also called a scent fingerprint or scent profile), which may be used to re-create and edit scents. Libraries of olfactory receptors are useful for determining the interaction pattern of a

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 8; Page 350-351; 1857pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English
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                                                                                                                                                                                                                                                                                                                                                                                                                                        3.2e-49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    s involved in olfactory
and antagonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           determination;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 948;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fuchs
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AAS15914;

AAS15914 standard;

CDNA;

25-JAN-2002

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CC The invention describes a novel isolated polypeptide, selected from a CC group of 21 G-protein coupled receptor polypeptides (GCREC) and useful in CC waccines and gene therapy. The polypeptide (I) is useful for screening CC for agonist or antagonist of (I), compounds specifically binding to (I), CC or compounds that modulate the activity of (I). The polymucleotide compounds that modulate the activity of (I). The polymucleotide in CC encoding (I) is useful for screening a compound for effectiveness in CC encoding (I) is useful for screening a compound for effectiveness in CC exposing a sample comprising the target polymucleotide to a compound, CC exposing a sample comprising the target polymucleotide, and comparing the expression of the target polymucleotide, and comparing CC disorders (e.g. cancers, arteriosclerosis, atherosclerosis), neurological disorders (e.g. epilepsy, stroke, schizophrenic disorders and compound (II) are CC disorders (e.g. epilepsy, stroke, schizophrenic disorders and comparing CC ischaemic heart disease), gastrointestinal disorders (e.g. anorexia, CC epetic ulcer), autoimmune/inflammatory disorders (e.g. diabetes mellitus, CC esteoporosis, psoriasis), and metabolic disorders such as obesity. CC esteoporosis, psoriasis), and metabolic disorders such as obesity. CC exposition profiles, to generate a transcript image of a tissue or cell expression profiles, to generate a transcript image of a tissue or cell type, and to generate hybridisation probes useful in mapping the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    anorexia; peptic ulcer; autoimmune disorder; inflammatory disor diabetes; osteoporosis; psoriasis; metabolic disorder; obesity; schizophrenic disorder; neuroskeletal disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cell proliferation disorder; cancer; arteriosclerosis; neurological disorder; epilepsy; stroke; cardiovascular disorder; hypertension; ischaemic heart disease; gastrointestinal disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  G-protein coupled receptor; GCREC; vaccine; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel G-protein coupled receptor polypeptides, for treating and preventing autoimmune/inflammatory disorders, neurological disorders, cell proliferative disorders, cardiovascular disorders and viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-656776/75.
P-PSDB; AAU10317.
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17-MAR-2000;
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naturally occurring genomic sequence. coupled receptor 18, one of 21 GCREC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DAM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5; Page 139-140; 141pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INCYTE GENOMICS INC
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e H, Khan FA,
Lo TP, Borows
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2000US-0190453P.
2000US-0190730P.
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/note= "G-protein coupled receptor 18"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  . 951
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          proteins
                                        This sequence encodes
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                described in
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, Yang J, Ha
                       the method
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Best Local
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24-MAR-2000; 2000US-0192033P.
12-APR-2000; 2000US-0198474P.
24-APR-2000; 2000US-019335P.
26-MAY-2000; 2000US-0207702P.
23-UUN-2000; 2000US-0213849P.
16-AUG-2000; 2000US-022534P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; olfactory receptor; G protein-coupled receptor; GPCR; odourant; ss; food additive; cosmetic; fragrance; pharmaceutical additive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human cDNA encoding olfactory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-MAR-2001; 2001WO-US007771.
                                                                                                                                                                                                                   Nucleic acids encoding human olfactory G protein-coupled receptors, useful for screening for compounds involved in olfactory sensation, the compounds can be used in the food, pharmaceutical and cosmetic industries to customize odors.
                                       olfactory sensation. The human olfactory receptors and polymucleotides encoding them are useful for screening a library of chemical compounds for compounds that are involved in olfactory sensation. Modulators of their activity are useful for pharmacological and genetic modulation of olfactory signalling pathways. Therefore, they can be used in the food, pharmaceutical and cosmetic industries to customise odours and
                                                                                                               The invention relates to nucleic acids encoding human olfactory receptors, OR, (a G protein-coupled receptor, GPCR). The OR's specifically recognise molecules, odourants, that elicit specific olfactory sensation. The human olfactory receptors and polynucleo
                                                                                                                                                                                                                                                                                           P-PSDB; AAU24614.
                                                                                                                                                                                      Claim 1; Page 128; 319pp; English.
                                                                                                                                                                                                                                                                                                                                                                 (SENO-)
                            fragrances.
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                                                                                                                                                                                                                                                                                                          2001-570867/64.
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2001US-0266862P.
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ative 0;
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Pred. No.
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                                 encodes a human olfactory receptor of
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Matches 120;
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ABZ43126
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                                                                                                                       for drug development e.g. based on olfactory and gustatory receptors in form of agonists and antagonists by screening intrinsic and extrinsic ligands as bitter taste inhibitors, taste enhancers and fragrance improvers. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directl from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                    Sequence 951 BP; 199
                                                                                                                                                                                        ABP95942) by extracting open-reading frames containing 6-8 transmembrane domains with 250-1000 amino acid residues to give a gene homologous with a known GPCR gene. The receptor proteins and encoded genes are useful for studying in vivo signal transduction mechanism and identifying targets for drug development e.g. based on olfactory and gustatory receptors in
                                                                                                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                                                                              Database global search for G protein-coupled receptors, proteins and encoded genes for studying in vivo signal transduction mechanism and identifying targets for drug development.
                                                                                                                                                                                                                                                          The invention relates to a method for screening G protein-coupled receptor (GPCR) genes (ABZ42870-ABZ43216) and/or GPCR proteins (ABP95596-
                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-304118/34.
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13-FEB-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NISC-)
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826 ACAAGACAACATCATCTCTGTTTTCTACACAATTGTCACTCCAGCCCTGAATCCACTCAT
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                                                              Similarity
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                                             Conservative (
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2001JP-00034434.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein coupled receptor; signal transduction;
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Pred. No.
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                                             Mismatches
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                                                                                                 G; 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             513.
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                                                       3.2e-49;
                                                                      DB 6;
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                                                                                                T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB
                                             0,
                                                                     Length 951
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                                          0;
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               885
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The invention relates to olfactory and Pheromone G-protein coupled CC receptor (GPCR) or a protein 95% identical to the GPCR, a specific active portion and its encoding polynucleotide. Also included are an agonist, antagonist or inhibitor of the GPCR or the polynucleotide a vector comprising the polynucleotide, a cell transformed by the vector, a non-the foliation of the polynucleotide comprising a partial or total deletion of the polynucleotide encoding the receptor and screening (detection and possibly, recovering) of compounds which are known or not known to be agonist, antagonists or compounds to the GPCR. The receptor-derived agonists, antagonists, or compounds to the GPCR. The receptor-derived comprovement, elimination or substitution of an existing taste and/or a fragrance of (or in) the food and/or cosmetic products. They can also be used in the preparation of medicament in the treatment and/or prevention of a mammalian disorder, such as cell migration, sterility, psychotic and depression, depression, for promoting axonal growth, nerve cell connection and nerve regeneration for modulating male and female connections and nerve regeneration of modulating male and female for the behaviours. Such as attimulation of several mammalian connections as attimulation of several mammalian.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 7
ABK68461
ID ABK6
behaviours, such as stimulation or suppression of appetite, sexual motivation, sexual attraction, aggression and for promoting or suppressing chemical communication between organisms. The present sequence is a human DNA encoding an olfactory and pheromone GPCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel pheromone G-protein coupled receptor and receptor-derived agonists, antagonists or inhibitors useful in food or cosmetic products or in the treatment or prevention of neurological disorders such as anxiety and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 196-197; 833pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            schizophrenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-330013/36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          aggression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CHEM-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              schizophrenia; manic depression; depression; axonal growth; menstrual cycle; appetite sexual motivation; sexual attraction;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ΒP
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Best Local S
Matches 120
                                                                              The invention relates to a method of representing sensory perception of CC one or more odourants. The method comprises: (a) providing a CC representative class of n olfactory receptors or ligand binding domains CC (LBD) of these receptors; (b) measuring values X1 to Xn representative of CC clast one activity of one or more odourants selected from: (i) binding CC one or more odourants to the LBD of at least one of the n olfactory receptors with CC receptors; (ii) activating at least one of the n olfactory receptors with the one or more odourants; and (iii) blocking at least one of the n of CC representation of sensory perception from the values X1 to Xn. The CC representation of the sensory perception of odourants is useful for the CC representation of foodurants of odourants is useful for the CC design and formulation of odourant human olfactory G-coupled receptor coding sequences and related PCR primers of the invention
Query Match
Best Local S
Matches 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA encoding G-coupled olfactory receptor #95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; olfactory G-coupled receptor; sensory perception of odourant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABK37593 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-JUN-2000; 2000US-0213812P.
13-MAR-2001; 2001US-00804291.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-DEC-2001.
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                                                                                                                                                                                                                                                                                                                          Representing sensory perception of one or more odorants for the identification and design of tastes and odors comprises providing representative group of n olfactory receptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-JUN-2001; 2001WO-US020122
                                                                                                                                                                                                                                                                                                 Example; Page 101; 182pp; English.
                                                        Sequence 951
                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SENO-)
                Local Similarity
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  120;
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                                                                                                                                                                                                                                                                                                                                                                                     AAU85234.
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 Conservative
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                                                         BP; 199
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                12.3%;
                                                         A; 275
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Pred. No.
    Score 120; DB b; -
; Pred. No. 3.2e-49;
                                                           C; 218
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                                                           G;
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                                                              U; 0 Other;
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08-FEB-2000; 2000US-0181004P.
08-FEB-2000; 2000US-0181013P.
08-FEB-2000; 2000US-0181013P.
08-FEB-2000; 2000US-0181043P.
24-UUL-2000; 2000US-0220262P.
25-UUL-2000; 2000US-02202694P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; olfactory receptor; OR; G-protein coupled receptor superfamily; GPCR; gene therapy; tranquiliser; vulnerary; anti-HTV; cytostatic; ds; nootropic; neuroprotective; antiparkinsonian; antiasthmatic; cancer; antiinflammatory; odorant receptor; olfactory loss; trauma; HTV; human immunodeficiency virus; neoplastic growth; neurological disorder; Parkinson's disease; Alzheimer's disease; neurogenesis; Crohn's disease; wound healing; asthma; Albright hereditary osteodystrophy; multiple sclerosis; nasal epithelial neuronal tissue; transgenic plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human odorant receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-FEB-2001; 2001WO-US003923.
                                                                                                                                                                                                           Novel isolated NOVX polypeptide related to human odorant receptor family of G-protein coupled receptor superfamily of proteins useful for treating trauma, neoplastic growth, Parkinson's disease, Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                       Padigaru M,
Baumgartner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-AUG-2000;
02-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CURA-)
                                                                                                                                                                                                                                                                                                                                                       2001-488883/53.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURAGEN CORP.
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), 2000US-0181043P.
), 2000US-022054P.
), 2000US-022054P.
), 2000US-022459P.
), 2000US-02452P.
                                                                                                                                                                                                                                                                                                                                                                                                            Spytek KA,
JC;
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2000US-0180646P.
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Novel isolated NOVX polynucleotides and their encoded polypeptides of t invention are related to the human olfactory (or odorant) receptors (ORs), from the G-protein coupled receptor (GPCR) superfamily. The sequences are useful for treating or preventing a pathology associated with OR in a subject, and for manufacturing a medicament for treating or preventing a pathology associated with OR in a subject, and for manufacturing a medicament for treating or preventing a medicament for treating and preventing and preventing a medicament for treating and preventing and preventing and preventing and preventing and preventing and preventing an

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Claim 9; Page 44; 199pp; English.

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RESULT 10
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Best Local Sim
Matches 120;
                                                                                                                                                                                                                                                            Alsobrook ;
Spytek KA;
                                                 New isolated olfactory receptor-like polypeptide, NOVX, useful for diagnosing, preventing or treating disorders of olfactory loss, e.g. trauma, human immunodeficiency virus illness, neoplastic growth and
                                               neurological
                                                                                                                                                                                                    WPI; 2002-471499/50.
                                                                                                                                                                                                                                                                                                                                                                                   02-NOV-2000; 2000US-0245292P.
06-FEB-2001; 2001US-00777789.
20-SEP-2001; 2001US-00245292.
                                                                                                                                                                                    P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NOV; odourant receptor; G protein coupled receptor; GPCR; trauma; HIV; ss; gene; olfactory receptor; olfactory loss; neoplastic growth; human immunodeficiency virus; Alzheimer's disease; neurological disord.
                                                                                                                                                                                                                                                                                                                                       (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            syndrome associated with a human disease. These diseases include disorders of olfactory loss, such as trauma, HIV illness and neoplastic growth, neurological disorders, such as Parkinson's disease and Alzheimer's disease, neurogenesis, cancer, wound healing, asthma, Crohn's disease, multiple sclerosis, and Albright hereditary osteodystrophy. The polynucleotides and polypeptides are also useful, to identify proteins of the same family, to screen for molecules which inhibit or enhance the neuronal tissue, in production of transgenic plants and for the development of new drug targets for various disorders. This sequence represents DNA encoding a human OR protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABK53096 standard;
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                                               disorders.
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Pred. No.
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Claim 9; Page 33; 122pp; English

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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC protein and antibody of the invention is useful for dispute. He way, compression and antibody of the invention is useful for disposing, which are disorders of olfactory loss, e.g. trauma, human immunodeficiency control (HIV) illness, neoplastic growth and neurological disorders, e.g. crambody is useful in screening assays, detection assays (e.g., contibody is useful in screening assays, detection assays (e.g., contibody is useful in screening assays, detection assays, predictive condicine (e.g., diagnostic assays, prognostic assays, monitoring clinical trials and pharmacogenomic), and in methods of treatment (e.g., condicine detection). The NOV protein is useful as immunogen to produce antibodies immunospecific for the protein, to screen for contential agonist and antagonist compounds, and as bait protein in a two-therapy, to express the protein, to detect NOVX mRNA or a genetic lesion invention is useful for isolating, and purifying the protein and to consider the protein in the protein in a sequence of the protein in the protein in a sequence of the construction is useful for isolating, and purifying the protein and to consider the protein in the protein in a sequence of the construction is useful for isolating, and purifying the protein and to consider the protein in the construction in the construc
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              09-OCT-2001; 2001WO-US031744
                                                                10-MAY-2002.
                                                                                                                                                                                                            variation
                                                                                                                                                                                                                                                                                 CDS
                                                                                                                                                                                                                                                                                                                                                                                NOV; odourant receptor; G protein coupled receptor; GPCR; trauma; HIV; ss; gene; olfactory receptor; olfactory loss; neoplastic growth; human immunodeficiency virus; Alzheimer's disease; SNP; neurological disorder; Parkinson's disease.
                                                                                                                                                                                                                                                                                                                                                  Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABK53097;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       useful for treating or preventing a disorder associated with NOV1 in subject, preferably human. A NOVX specific antibody is useful for determining the presence or amount of protein in a sample. The DNA, protein and antibody of the invention is useful for diagnosing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to the DNA and protein sequences isolated olfactory receptor-like polypeptide, NOVX. The sequences of the invention and an antibody specific for useful for treating or preventing a disorder associated.
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Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     encoding novel odourant receptor NOV7 protein.
                                                                                                                                                                                     /*tag= a
/product= "NOV7 protein"
replace(278,T)
                                                                                                                                                                                                                                                                            Location/Qualifiers 27. .1001
                                                                                                                                        standard_name= "Single nucleotide polymorphism"
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This invention relates to the DNA and protein sequences of a novel isolated olfactory receptor-like polypeptide, NOVX. The DNA and protein are sequences of the invention and an antibody specific for the protein are consequences of the invention and an antibody specific for the protein are subject, preferably human. A NOVX specific antibody is useful for consider associated with NOVI in a compount of protein in a sample. The DNA, consider and antibody of the invention is useful for disquosing, consider as disorders of olfactory loss, e.g. trauma, human immunodeficiency virus (HIV) illness, neoplastic growth and neurological disorders, e.g. virus (HIV) illness, neoplastic growth and neurological disorders, e.g. parkinson's disease and Alzheimer's disease. The sequences and NOV consomal mapping, tissue typing, forensic biology), predictive chromosomal mapping, tissue typing, forensic biology), predictive chromosom
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Best Local Similarity
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06-FEB-2001; 2001US-00777789
20-SEP-2001; 2001US-00245292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated olfactory receptor-like polypeptide, NOVX, useful for diagnosing, preventing or treating disorders of olfactory loss, e.g. trauma, human immunodeficiency virus illness, neoplastic growth and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1008 BP;
                                                                                                                                                                                                                        AAS11687 standard; DNA; 1050
Human; olfactory receptor; OR; G-protein coupled receptor superfamily; GPCR; gene therapy; tranquiliser; vulnerary; anti-HIV; cytostatic; ds; nootropic; neuroprotective; antiparkinsonian; antiasthmatic; cancer;
                                                                                      Human odorant receptor (OR)
                                                                                                                                      24-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2002-471499/50
                                                                                                                                                                                                                                                                                                                                        917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                120;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU97928.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACAAGACAACATCATCTCTGTTTTCTACACAATTGTCACTCCAGCCCTGAATCCACTCAT 885
                                                                                                                                                                                                                                                                                                                                                                          CTACAGCCTGAGGAATAAGGAGGTCATGCGGGCCTTGAGGAGGGTCCTGGGAAAATACAT
                                                                                                                                                                                                                                                                                                                                                                                                                          acaagacaacarcarcrererreracacaarrereacciccaecereaarceacrecar
                                                                                                                                                                                                                                                                                                                                     cTACAGCCTGAGGAATAAGGAGGTCATGCGGGCCTTGAGGAGGGTCCTGGGAAAATACAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Burgess
                                                                                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  216 A; 290 C; 234 G; 268 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12.3%; 50.
100.0%; Pr
                                                                                                                                        entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 120;
pred. No.
                                                                                                 DNA #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lepley DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.2e-49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 1008;
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ACAAGACAACATCATCTCTGTTTTCTACACAATTGTCACTCCAGCCCTGAATCCACTCAT

Matches Query Match

Local Similarity

Conservative

0

Mismatches

12.3%;

Score 120; Pred. No.

3.2e-49; DB T; 0 5

Indels

0

Gaps

885 0 Length 1050; U; 0 Other;

Sequence

1050

BP;

234 A; 293 C; 238 G; 285

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antiinflammatory; odorant receptor; olfactory loss; trauma; HIV; human immunodeficiency virus; neoplastic growth; neurological disorder; parkinson's disease; Alzheimer's disease; neurogenesis; Crohn's disease wound healing; asthma; Albright hereditary osteodystrophy; multiple sclerosis; nasal epithelial neuronal tissue; transgenic plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-FEB-2000;
07-FEB-2000;
07-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-AUG-2001.
                                                                                                  Novel isolated NOVX polynucleotides and their encoded polypeptides of the invention are related to the human olfactory (or odorant) receptors (OR8), from the G-protein coupled receptor (GPCR) superfamily. The sequences are useful for treating or preventing a pathology associated with OR in a subject, and for manufacturing a medicament for treating a with OR in a subject, and for manufacturing a medicament for treating a syndrome associated with a human disease. These diseases include syndrome associated with a human disease. These diseases and neoplastic disorders of olfactory loss, such as trauma, HIV illness and neoplastic growth, neurological disorders, such as Parkinson's disease and Albriemer's disease, neurogenesis, cancer, wound healing, asthma, Crohn's Alzheimer's disease, neurogenesis, and Albright hereditary osteodystrophy. The disease, multiple sclerosis, and Albright hereditary osteodystrophy. The polymucleotides and polypeptides are also useful, to identify proteins of the same family, to screen for molecules which inhibit or enhance the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-FEB-2000;
08-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-FEB-2001; 2001WO-US003923
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-FEB-2000;
24-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                         Novel isolated NOVX polypeptide related to human odorant receptor family of G-protein coupled receptor superfamily of proteins useful for treating trauma, neoplastic growth, Parkinson's disease, Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Baumgartner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Padigaru M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-AUG-2000;
                                                                                                                                                                                                                                                                                                                                           Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                 P-PSDB; AAU07087.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CURA-)
                                 activity or function of olfactory receptors, to detect nasal epithelial neuronal tissue, in production of transgenic plants and for the development of new drug targets for various disorders. This sequence represents DNA encoding a human OR protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001-488883/53.
                                                                                                                                                                                                                                                                                                                                           9; Page 39; 199pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000US-0181043P.
2000US-0220262P.
2000US-0220594P.
2000US-0224596P.
2000US-0224592P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         spytek KA,
JC;
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2000US-0180646P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000US-0181013P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001US-00777789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-0181004P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers 72. .1022
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /product= "Human odorant receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ballinger RA, Mishra VS;
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RESULT 13
ADC85922
ID ADC85
XX ADC85922
XX ADC85
         RESULT 14
AAA51192
ID AAA51
XX
AC AAA51
XX
DT 26-SE
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Best Local Simi
Matches 120;
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       26-SEP-2000
                                                      AAA51192;
                                                                                                  AAA51192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel polynucleotide encoding a guanosine triphosphate-binding protein coupled receptor (GPCR). A polynucleotide the invention may have a use in gene therapy. The polynucleotide and polypeptide are useful for preparing a composition for treating a patie in need of increased or suppressed activity or expression of the guanosine triphosphate-binding protein coupled receptor. The polynucleotide sequences shown in ADC85548-ADC87816 encode GPCR's of the immeria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID NO 275; 28pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polynucleotide, useful for preparing a composition for treating a patient in need of increased or suppressed activity or expression of the guanosine triphosphate-binding protein coupled receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Suwa M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-JUN-2001; 2001JP-00246789.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               guanosine triphosphate-binding protein coupled receptor; gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADC85822 standard; DNA; 1351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ds; gene; human; GPCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NAAD-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC85822;
                                                                                                                                                                                                                                                                                                              1007
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                                                                                                                                                                                                                                                                  886
                                                                                                                                                                                                                                                                                                                        826 ACAAGACAACATCATCTCTGTTTTCTACACAATTGTGACTCCAGCCCTGAATGCACTCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2003-315783/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        988
                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAT INST ADVANCED IND SCI & TECHNOLOGY.
CENT ADVANCED SCI & TECHNOLOGY INCUBATIO.
                                                                                             standard;
                                                                                                                                                                                               CTACAGCCTGAGGAATAAGGAGGTCATGCGGGCCTTGAGGAGGGTCCTGGGAAAATACAT
                                                                                                                                                                                                                             CTACAGCCTGAGGAATAAGGAGGTCATGCGGGCCTTGAGGAGGGTCCTGGGAAAATACAT
                                                                                                                                                                                                                                                                                             ACAAGACAACATCATCTGTTTTCTACACAATTGTCACTCCAGCCCTGAATCCACTCAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Asai K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTACAGCCTGAGGAATAAGGAGGTCATGGGGGCCTTGAGGAGGGTCCTGGGAAAATACAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTACAGCCTGAGGAATAAGGAGGTCATGCGGGCCTTGAGGAGGTCCTGGGAAAATACAT
                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 342 A; 340 C;
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                                                                                             CDNA; 678
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                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                        Score 120;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Aburatani H;
                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          295 G; 374 T;
                                                                                                                                                                                                                                                                                                                                                                                                           3.2e-49;
                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 1351;
                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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ACA0491
                                                                                                                                                                                                                                                                                                                    RESULT 15
                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 27
ss; olfactory receptor; odourant; ligand-binding region; mouse; olfactory neuron; olfactory epithelium; 7 transmembrane recepto olfactory response.
                                                                                                                                                                                                                                                                                                                                         AAA51175-93 encode murine odorant/ligand binding regions of olfactory receptors. They were amplified using degenerate primers shown in AAA51163 for the primers can be used to generate a library of OR-LBR comprising the transmembrane domains (TMD) II-VI, III-VI, II-VII or III-VII. Chimeric nucleic acids encoding a 7-TMD protein comprise a sequence encoding an N-terminal plasma membrane translocation domain (especially translocation domain was initially derived from the N-LBR. The human rhodopsin receptor. LBR produced by amplification with the primers are useful for generating new odorants, to screen for toxicity or therapseutic activity in odorants, and altering an animal's olfactory
                                                     Mouse odourant/ligand binding region cDNA #18.
                                                                                       28-MAY-2003
                                                                                                                  ACA04914;
                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                        ACA04914 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 18; Page 36-37; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primer pair for amplifying olfactory receptor nucleic acid, useful e.g. for producing receptor libraries used in e.g. screening odorants for toxicity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Degenerate primer; ligand-binding region; olfactory receptor; Transmembrane; translocation human rhodopsin receptor; odorant; toxicity; olfactory response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Murine olfactory receptor ligand-binding region cDNA 18
                                                                                                                                                                                                                      370 CATGGCCTATGACAGGTATGTGGCCAT 396
                                                                                                                                                                                                             162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000-431471/37.
                                                                                                                                                                                                                                                               l Similarity
27; Conserv
                                                                                                                                                                                                    CATGGCCTATGACAGGTATGTGGCCAT 188
                                                                                                                                                                                                                                                                                                              678 BP; 148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                UNIV JOHNS HOPKINS SCHOOL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Krautwurst
                                                                                                                                                                                                                                                            Conservative
                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98US-00112605.
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/product=
/partial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                        cDNA; 678
                                                                                                                                                                                                                                                                                                          A; 174 C; 119 G; 237 T; 0 U;
                                                                                                                                                                                                                                                                         100.0%;
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                                                                                                                                                                                                                                                                                       2.8%;
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                                                                                                                                                                                                                                                            0;
                                                                                                                                        ΒP
                                                                                                                                                                                                                                                                       Score 27;
Pred. No.
                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDICINE
                                                                                                                                                                                                                                                    DB 3; LC
3. 0.0069;
0;
             transmembrane receptor;
                                                                                                                                                                                                                                                                                Length 678;
                                                                                                                                                                                                                                                                                                          0 Other;
                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                     Gaps
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Mus musculus.

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Page 10
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Search completed: October 6, 2004, 03:52:22
Job time : 436 secs
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                                                                                                                                        Query Match 2.8%; Score 27; DB 7; Louist Local Similarity 100.0%; Pred. No. 0.0069; Matches 27; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                        The invention relates to an amplification primer sequence pair for amplifying a nucleic acid encoding an olfactory receptor ligand-binding region, comprising a primers appearing as ACA04888 and ACA04889. Also included is a kit for amplifying olfactory receptor sequences comprising the primers. The primers are useful for generating a nucleic acid sequence that encodes a ligand-binding region of an olfactory receptor. The amplified nucleic acid is genomic DNA, mank or cDNA derived from olfactory neurons or the olfactory epithelium. The olfactory receptors are 7 transmembrane receptors, manipulation of which can modulate an animals olfactory response and can be used to generate novel odourants. The present sequence encodes a mouse olfactory receptor ligand binding region and is isolated using the primer of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US6492143-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel amplification primer sequence pair for amplifying nucleic acid encoding an olfactory receptor ligand-binding region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYJO ) UNIV JOHNS HOPKINS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Col 77-80; 46pp; English.
                                                                                                                                                                                                             Sequence 678 BP; 148 A; 174 C; 119 G; 237 T; 0 U; 0 Other;
                                                                                        370 CATGGCCTATGACAGGTATGTGGCCAT 396
                                                                      162 CATGGCCTATGACAGGTATGTGGCCAT 188
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US-09-465-901-45
                                                                                                                                                                  RESULT 1
PATENT No. 6492143
GENERAL INFORMATION:
APPLICANT: Reed, Randall
APPLICANT: Yeau, King-Wai
APPLICANT: Krautwurst, Dietmar
TITLE OF INVENTION: Olfactory Receptor Expession Libraries
TITLE OF INVENTION: ad Methods of Making and Using Them
FILE REFERENCE: 001107.00105
CURRENT APPLICATION NUMBER: US/09/465,901
CURRENT FILING DATE: 2001-12-17
                                                                                                                                                                                                                                                                                                                                                                                      Result
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Database
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Maximum
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                                                                                                                 Scoring table:
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DB
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seq length:
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                                                                                                                                     Application US/09465901
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Match Length
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                                                                                                                                                                                                                                              /cgn2 6/ptodata/2/ina/5A_COMB.seq:*
/cgn2 6/ptodata/2/ina/5B_COMB.seq:*
/cgn2 6/ptodata/2/ina/6A_COMB.seq:*
/cgn2 6/ptodata/2/ina/6B_COMB.seq:*
/cgn2 6/ptodata/2/ina/6B_COMB.seq:*
/cgn2 6/ptodata/2/ina/PCTUS_COMB.seq:*
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US-09-465-901-47

US-08-599-252-84

US-08-436-074-57

PCT-US96-06352-84

PCT-US96-06583-84
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US-08-884-324-13
US-08-884-324-14
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(without alignments)
5315.559 Million cell updates/sec
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APPLICANT: Yau, King-Wai
APPLICANT: Yau, King-Wai
APPLICANT: Yau, King-Wai
APPLICANT: Krautwurst, Dietmar
TITLE OF INVENTION: Olfactory Receptor Expession Libraries
TITLE OF INVENTION: ad Methods of Making and Using Them
FILE REFERENCE: 001107.00105
CURRENT APPLICATION NUMBER: US/09/465,901
CURRENT FILING DATE: 2001-12-17
PRIOR APPLICATION NUMBER: 60/112,605
PRIOR APPLICATION NUMBER: 60/112,605
PRIOR FILING DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 57
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 47
LENGTH: 951
TYPE: DNA
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                                                                                                                                                                                                                                Sequence 84, Application US/08599252
Patent No. 5705343
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local (
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Patent No. 6492143
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PRIOR FILLING DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 57
SOFTWARE: FAStSEQ for Windows Version
SEQ ID NO 45
LENGTH: 678
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                            APPLICANT: GNIRKE, ANDREAS
APPLICANT: KIMMEL, BRUCE E.
APPLICANT: THOMAS, WINSTON J.
APPLICANT: WOLFF, ROGER K.
TITLE OF INVENTION: METHOD TO DIAGNOSE HEREDITARY
TITLE OF INVENTION: HENOCHROMATOSIS
NUMBER OF SEQUENCES: 124
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
                                                                                                                                                          APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Mus musculus FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
                ADDRESSEE: MORRISON & FOER STREET: 2000 Pennsylvania
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27; Conserva
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2.3%; Score 22;
Similarity 100.0%; Pred. No.
22; Conservative 0; Mismatc
Washington
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Conservative 0; Mismatches
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Pred. No.
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STATE: USA COUNTRY: USA 20006-1888 TE: 2000ABLE F

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US-08-436-074-57/c
; Sequence 57, Application US/08436074
; Patent No. 5753438
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/599,252
FILING DATE: 09-FEB-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 9053-0001.21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 22; Conserv
                                                                                                                                                                                                                                                                                 COUNTRY: USA
COUNTRY: USA
ZIP: 20006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/436,074
FILING DATE: US-MAY-1995
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LENGTH: 1320 base pair
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TELEPHONE: (202) 887-1500
TELEFAX: (202) 887-0763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: GNIRKE, ANDREAS
APPLICANT: KIMMEL, BRUCE E.
APPLICANT: THOMAS, WINSTON J.
APPLICANT: WOLFF, ROGER K.
TITLE OF INVENTION: METHOD TO DIAGNOSE HEREDITARY
TITLE OF INVENTION: HEMOCHROMATOSIS
                          CLASSIFICATION: 436
ATTORNEY/AGENT INFORMATION:
NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 9053-0001.00
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000 Pennsylvania Ave. N.W., Suite 5500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEDER, JOHN N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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(202) 887-1500
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Pred. No. 0.22;
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PCT-US96-06352-84/c
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Best Local Similarity 100.0%;
Chas 22; Conservative (
                                                             ; TOPOLOGY:
PCT-US96-06352-84
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1320 base pair
Query Match
Best Local S
Matches 22
                                                                                                                                   TELEX: 90-4030
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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APPLICANT:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ADDRESSEE: MORRISON & FOERSTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: WOLFF, ROGER K.
TITLE OF INVENTION: METHOD TO DIAGNOSE HEREDITARY
TITLE OF INVENTION: HEMOCHROMATOSIS
                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: PCT/US96/06352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08 FILING DATE: 09-FEB-1996 ATTORNEY/AGENT INFORMATION: NAME: MURASHIGE, KATE H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                    REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 9053-0001.21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-1500
TELEFAX: (202) 887-0763
                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 1320 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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ZIP: 20006-1888
                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                            STRANDEDNESS:
                                                                                                                        LENGTH:
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il Similarity
22; Conserva
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GNIRKE, ANDREAS
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 2.3%; Score 22; ilarity 100.0%; Pred. No. Conservative 0; Mismatc
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Pred. No.
      0; Mismatches
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                      DB 5;
0.22;
                                     Length 1320;
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846 TTTTCTACACAATTGTCACTCC 867

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1210 TTTTCTACACAATTGTCACTCC 1189

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PCT-US96-06583-84/c
                                                                                                                                                            Sequence 12, Application US/08884324 Patent No. 6060283
                                                                                                                                                                                                                                                                                                                                        Best Local Similarity Matches 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                               GENERAL INFORMATION:
APPLICANT: Takanori OKURA
APPLICANT: Kakuji TONIGOE
APPLICANT: Masahi KURIMOTO
TITLE OF INVENTION: GENOMIC DNA
TITLE OF INVENTION: OF INDUCING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/599,252

PILING DATE: 09-FEB-1996

ATTORNEY/AGENT INFORMATION:

NAME: MURASHIGE, KATE H.

REGISTRATION NUMBER: 29,959

REFERENCE/DOCKET NUMBER: 9053-0001.21

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: 90-4030
INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 1320 base pairs
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CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK
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TITLE OF INVENTION: HEM
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 20006-1888

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Re-lease #1.0, Version #1.30
CURRENT APPLICATION DATA:
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TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
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TELEFAX: (202) 887-0763
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CITY: Washington
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THOMAS, WINSTON J.
WOLFF, ROGER K.
NVENTION: METHOD TO DIAGNOSE HEREDITARY
VYENTION: HEMOCHROMATOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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FEDER, JOHN N.
GNIRKE, ANDREAS
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100.0%; Pred. No.
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                                             ENCODING A POLYPEPTIDE CAPABLE THE PRODUCTION OF INTERFERON-
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PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: JP 185,305/96
APPLICATION NUMBER: 27-JUN-1996
ATTORNEY/AGENT INFORMATION:
NAME: BROWDY, Roger L.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: OKURA=1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 202-737-3528
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Kakuji TORIGOE
APPLICANT: Masahi KURIMOTO
TITLE OF INVENTION: GENOMIC DNA ENCODING A POLYPEPTIDE CAPABLE
TITLE OF INVENTION: OF INDUCING THE PRODUCTION OF INTERFERON-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: (
ORIGINAL SOURCE:
ORGANISM: huma
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/884,324
                                                                                                                                                                                                                                                                                                   STREET:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                 STATE:
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LOCATION:
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mes 20; Conserv
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TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                831 ACAACATCATCTCTGTTTTC 850
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APPLICATION DATA:
                                                                                                                                                                                                20004
                                                                                                                                                                                                                                                                                     Washington
                                                                                                                                                                                                                                                             D.C.
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                                                                                                                                                                                                                                                                                                                    E: BROWDY AND NEIMARK
419 Seventh Street, N.W.,
                                                                                                                                                                                                                              USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Takanori OKURA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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rive 0; Mismatches
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                        Version #1.30
                                                                                                                                                                                                                                                                                                                        Suite 300
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US-08-884-324-13
                                                         Query Match 2.0%;
Best Local Similarity 100.0%;
Matches 20; Conservative
                                                        Matches
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SEQUENCE CHARACTERISTICS:
LENGTH: 11464 base pairs
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REFERENCE/DOCKET NUMBER: OX
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEPAX: 202-737-3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: JP 185,305/96 FILING DATE: 27-JUN-1996 ATTORNEY/AGENT INFORMATION:
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ORIGINAL SOURCE:
ORGANISM: human
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NAME/KEY: leader
LOCATION
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REGISTRATION NUMBER: :
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LOCATION: 1..3
IDENTIFICATION METHOD:
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[DENTIFICATION METHOD:
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LOCATION: 11444..11464
IDENTIFICATION METHOD:
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4226 ACAACATCATCTCTGTTTTC 4207
                  831 ACAACATCATCTCTGTTTTC 850
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4866..4983
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Pred. No. 2.3;
                                                           Mismatches
                                                              0; Indels
                                                                                         Length 11464;
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US-08-884-324-14/c

RESULT 9

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Sequence 14, Application US/08884324 patent No. 6060283
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 28994 base pai
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APPLICANT:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: KAKUJI TORIGOE
APPLICANT: MASSAH KURIMOTO
TITLE OF INVENTION: GENOMIC DNA ENCODING A POLYPEPTIDE CAPABLE
TITLE OF INVENTION: OF INDUCING THE PRODUCTION OF INTERFERON-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 10
APPLICATION UNMBER: JP 10
FILING DATE: 27-JUN-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: BROWDY, ROGER L.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: OK
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
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ORIGINAL SOURCE:
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FILING DATE:
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               NAME/KEY: intron
LOCATION: 20587..21920
IDENTIFICATION METHOD: E
NAME/KEY: mat peptide
LOCATION: 21921..22054
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NAME/KEY: mat peptide
LOCATION: 20469..20586
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419 Seventh Street, N.W., Suite 300
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20452..20468
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NAME/KEY: intron

LOCATION: 22055..26827

IDENTIFICATION METHOD: E

NAME/KEY: mat peptide

LOCATION: 26828..27046

IDENTIFICATION METHOD: S

NAME/KEY: 3'UTR

LOCATION: 27047..28994

IDENTIFICATION METHOD: E

US-08-884-324-14

Query Match
Best Local Similarity 100.0%; Pred. No. 2.4;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Matches 20; Conservative 101.0% pred. No. 2.4;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy

831 ACAACATCATCTCTGTTTTC 850

Db 19829 ACAACATCATCTCTGTTTTC 19810

Search completed: October 6, 2004, 05:40:52

Job time: 102 secs
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Result
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AAU24616
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(without alignments)
731.844 Million cell updates/sec
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	Human	Human	Human	Human	_	Aag72799 Human olf	Aag72749 Human olf	Aar27879 Pheromone	_	Human	Aag71547 Human olf	Ade08306 Novel pro	Adc86715 Human GPC	Adc86807 Human GPC		Aab45704 Human 7TM	Adc86213 Human GPC	Aau85236 G-coupled	Aau95552 Human olf	

ALIGNMENTS

AAG71681 RESULT 1 Human; olfactory receptor; OR; primary scent determination; secondary scent determination; polypeptide library; odour receptor; scent profile; scent fingerprint; scent representation. 08-OCT-1999; 99US-0158615P. 24-FEB-2000; 2000US-0184809P. 06-OCT-2000; 2000WO-US027582 19-APR-2001. WO200127158-A2 Homo sapiens. Human olfactory receptor polypeptide, 30-JUL-2001 AAG71681; AAG71681 standard; protein; 316 (first entry) AA SEQ ID NO:

(YEDA) YEDA RES & DEV CO LTD.

Bellenson J, Smith D, Lancet D, Glusman G, Fuchs T, Yanai I;

WPI; 2001-290713/30.

New polynuclectides which encode polypeptides involved in olfactory sensation for identifying olfactory agonists and antagonists.

Claim 11; Page 845; 1857pp; English.

The present sequence is an olfactory receptor which is encoded by one of a number of novel polynucleotides. The polynucleotides can be used in screening for olfactory agonists and antagonists. The methods allow for the determination of primary scents and the identification of the odour receptors used to detect these primary scents. The methods also enable determination of secondary scents and the identification of combinations of odour receptors that are involved in detecting such secondary scents. This enables the construction of a scent representation (also called a scent fingerprint or scent profile), which may be used to re-create and edit scents. Libraries of olfactory receptors are useful for determining the interaction pattern of a composition with the receptors, and can be

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                                  Padigaru M,
Conley P, H
Edinger S,
                                                                                                                                                                                                                                                                                 GPCR1; olfactory receptor; G-protein coupled receptor; receptor; human; antiarteriosclerotic; antidiabetic; analgesic; cytostatic; antidepressant; nootropic; neuroprotective; immunosuppressive; antiallergic; antibacterial; virucide; protozoacide; fungicide; cardiomyopathy; atherosclerosis; diabetes; gene therapy.
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                                                                                                                                                                                                                                                                                                                                                         Human olfactory receptor-like protein GPCR1.
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                                                                                                                                                                                                                                                               Homo sapiens.
  N-PSDB; ABQ81238
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             WPI; 2002-698753/75
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                                             , Gerlach VL,
Hart M, Tomli
                                                                                  THERAPEUTICS
                                      Macdougall
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                                                 Tomlinson JE,
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Kekuda R,
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XEFEEXXXX55555555555555555555555555555
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Claim 1; Page 11; 111pp; English.

transmembrane receptor (rhodopsin family). The invention provides GPCR1 polypeptides and nucleic acids, and methods for their recombinant provides and nucleic acids, and methods for their recombinant production. These are useful for treating or preventing a GPCR1-associated disorder, such as cardiomyopathy, atherosclerosis, diabetes, or a disorder related to cell signal processing or metabolic pathway or modulation in humans (claimed). They may further be used for treating or preventing e.g. developmental diseases, metabolic pathway disorders, retinal disorders, wasting disorders associated with chronic diseases, retinal disorders, wasting disorders associated with chronic diseases, pain, cancer, psychotic and neurological disorders (e.g. anxiety or schizophrenia), autoimmune diseases, allergies, bacterial, fungal, protozoal and viral infections, neurodegenerative diseases (e.g. nucleic acids may also be used as research tools, as diagnostic or nucleic acids may also be used as research tools, as diagnostic or in the control of the control The present sequence is the protein sequence of a novel human G-protein coupled receptor, designated GPCR1, that has structural and physiological functions characteristic of the olfactory receptor family, and has properties similar to those of proteins containing the 7tm 1 7 prognostic markers, in gene therapy, for promoting tissue regeneration vitro and in vivo, and as biological defence weapons. The proteins are also are useful for producing antibodies specific for the proteins, as vaccines, and in screening for potential agonists and antagonists ij

Sequence 316 AA;

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Similarity 100.0%; Pred. No. 1.2e-281;
16; Conservative 0; Mismatches 0;
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                                                         CSSHLIVVGMFYGAATFMYVLPSSFHSPKQDNIISVFYTIVTPALNPLIYSLRNKEVMRA
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RESULT 3
AAG71678
ID AAG7
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AC AAG7
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DT 30-J
XX
DE Huma
XX
KW Huma
KW Becc
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                                        30-JUL-2001
                                                                        standard;
                                         (first entry)
                                                                        protein; 316
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Human olfactory receptor polypeptide, SEQ ID NO: 1359

Human; olfactory secondary scent / receptor; OR;
determination; primary scent determination;
polypeptide library; odour receptor;

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RESULT 4
AAU10317
ID AAU1
XX AAU1
XX AAU1
XX AAU1
XX AC
DT 25-J
XX G-pr
XX G-pr
XX G-pr
XX Cell
XW neur
KW hype
KW hype
KW diab
XW Schi
XW Geh
XW Geh
XW Geh
XW Homo
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                                                                                  neurological disorder; epilepsy; stroke; cardiovascular disorder; hypertension; ischaemic heart disease; gastrointestinal disorder; anorexia, peptic ulcer; autoimmune disorder; inflammatory disorder; diabetes; osteoporosis; psoriasis; metabolic disorder; obesity; schizophrenic disorder; neuroskeletal disorder.
                                                                                                                                                             G-protein coupled receptor; GCRBC; vaccine; gene therapy; cell proliferation disorder; cancer; arteriosclerosis;
                             WO200166742-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           determination of secondary scents and the identification of combinations of odour receptors that are involved in detecting such secondary scents. This enables the construction of a scent representation (also called a scent fingerprint or scent profile), which may be used to re-create and edit scents. Libraries of olfactory receptors are useful for determining the interaction pattern of a composition with the receptors, and can be used for determining differences in the olfactory faculties of different
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is an olfactory receptor which is encoded by one of a number of novel polymucleotides. The polymucleotides can be used in screening for olfactory agonists and antagonists. The methods allow for the determination of primary scents and the identification of the odour receptors used to detect these primary scents. The methods also enable
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      individuals
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57; Conserv
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YEDA RES &
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                                                                                                                                                                                                                                                                                                            standard; protein; 316 AA
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2000US-0184809P.
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and antagonists.
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                                                                                                                                                                                                                                                                                                                                                                                     The invention describes a novel isolated polypeptide, selected from a C group of 21 G-protein coupled receptor polypeptides (GCREC) and useful in CC vaccines and gene therapy. The polypeptide (I) is useful for screening CC for agonist or antagonist of (I), compounds specifically binding to (I) corresponds that modulate the activity of (I). The polymucleotide corresponds that modulate the activity of (I). The polymucleotide compounds specifically binding to (I) corresponds that modulate the activity of (I). The polymucleotide compound for effectiveness in CC altering expression of a target polymucleotide comprising (II), by CC detecting asmple comprising the target polymucleotide to a compound, CC detecting altered expression of the target polymucleotide, and comparing CC exposing a sample comprising the target polymucleotide, and comparing CC disorders (e.g. cancers, arteriosclerosis, atherosclerosis), neurological CC disorders (e.g. cancers, arteriosclerosis, atherosclerosis), neurological CC disorders (e.g. epilepsy, stroke, schizophrenic disorders (e.g. anorexia, CC peptic ulcer), autoimmune/inflammatory disorders (e.g. hypertension, CC disorders (e.g. autoimmune/inflammatory disorders (e.g. diabetes mellitus, CC peptic ulcer), autoimmune/inflammatory disorders such as obesity. CC peptic ulcer) autoimmune/inflammatory disorders such as obesity. CC peptic ulcer) polymucleotide is useful; as primers for detecting costeoporosis, psoriasis), and metabolic disorders such as obesity. The polymucleotide is useful in mapping the comparate hybridisation probes useful in mapping the maturally occurring genomic sequence. This is the G-protein coupled cof the invention
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Best Local :
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17-MAR-2000;
20-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel G-protein coupled receptor polypeptides, for treating preventing autoimmune/inflammatory disorders, neurological cell proliferative disorders, cardiovascular disorders and very cell proliferative disorders.
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; 2000US-0190453P.
; 2000US-0190730P.
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Khan FA, Policky JL,
                                                                                                                                 protein;
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                                                                                                                                                                                                                                                                                                  Score 57;
Pred. No.
                                                                                                                                                                                                                                                                              Mismatches
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Au-Young J,
                                                                                                                                                                                                                                                                                                              Length 316;
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                                                                                                                                                                                                                                                                              Gaps
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Human; olfactory receptor; G protein-coupled receptor; GPCR; odourant;

Human olfactory receptor AOLFR107.

18-DEC-2001

(first entry)

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RESULT 6
AAU07087
ID AAU0
XX AAU0
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XX Huma
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24-MAR-2000;
12-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to nucleic acids encoding human olfactory receptors, OR, (a G protein-coupled receptor, GPCR). The OR's specifically recognise molecules, odourants, that elicit specific olfactory sensation. The human olfactory receptors and polynucleotides encoding them are useful for screening a library of chemical compounds for compounds that are involved in olfactory sensation. Modulators of their activity are useful for pharmacological and genetic modulation of olfactory signalling pathways. Therefore, they can be used in the food, pharmaceutical and cosmetic industries to customise odours and fragrances. The present sequence is a human olfactory receptor of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200168805-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acids encoding human olfactory G protein-coupled receptors, useful for screening for compounds involved in olfactory sensation, the compounds can be used in the food, pharmaceutical and cosmetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             food additive; cosmetic; fragrance; pharmaceutical additive
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07-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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nootropic; neuroprotective; antiparkinsonian; antiasthmatic; cancer; antiinflammatory; odorant receptor; olfactory loss; trauma; HIV; human immunodeficiency virus; neoplastic growth; neurological disorder; parkinson's disease; Alzheimer's disease; neurogenesis; Crohn's disease;
                                                                          Human; olfactory receptor; OR; G-protein coupled receptor superfamily; GPCR; gene therapy; tranquiliser; vulnerary; anti-HIV; cytostatic;
                                                                                                                                          Human odorant
                                                                                                                                                                                                                                                               AAU07087 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 industries to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SENO-) SENOMYX INC
                                                                                                                                                                                                                       AAU07087;
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)B; AAS42307.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                           BARLHMPMYLLLGQLSIMDLLFTSVVTPKALADFIRRENTISFGGCALQMFLALIMG 108
                                                                                                                                                                                                                                                                                                                                                                EARLHMPMYLLLGQLSLMDLLFTSVVTPKALADFLRRENTISFGGCALQMFLALTMG 108
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2000US-0207702P.
2000US-0213849P.
2000US-0226534P.
2000US-0230732P.
2001US-0266862P.
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2000US-0198474P.
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Pred. No.
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ABP95852
ID ABP9
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AC ABP9
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DT 06-M
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06-MAR-2003 ABP95852; ABP95852

(first

entry)

standard;

protein; 316

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52

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Novel isolated NOVX polynucleotides and their encoded polypeptides of the cinvention are related to the human olfactory (or odorant) receptors (CC (ORs), from the G-protein coupled receptor (GPCR) superfamily. The CC (ORs), from the G-protein coupled receptor (GPCR) superfamily. The CC with OR in a subject, and for manufacturing a pathology associated CC with OR in a subject, and for manufacturing a medicament for treating a cc syndrome associated with a human disease. These diseases include cc disorders of olfactory loss, such as trauma, HIV illness and neoplastic CC growth, neurological disorders, such as trauma, HIV illness and neoplastic CC disease, meurogenesis, cancer, wound healing, asthma, Crohn's CC disease, multiple sclerosis, and Albright hereditary osteodystrophy. The CC polynucleotides and polypeptides are also useful, to identify proteins of the same family, to screen for molecules which inhibit or enhance the CC activity or function of olfactory receptors, to detect nasal epithelial neuronal tissue, in production of transgenic plants and for the CC development of new drug targets for various disorders. This sequence
                                                                          Matches
                                                                                                  Query Match
Best Local
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08-FEB-2000;
08-FEB-2000;
08-FEB-2000;
08-FEB-2000;
08-FEB-2000;
24-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Padigaru M,
Baumgartner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-JUL-2000;
11-AUG-2000;
02-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          wound healing; asthma; Albright hereditary osteodystrophy; multiple sclerosis; nasal epithelial neuronal tissue; transgenic plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel isolated NOVX polypeptide related to human odorant receptor family of G-protein coupled receptor superfamily of proteins useful for treating trauma, neoplastic growth, Parkinson's disease, Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-FEB-2000;
07-FEB-2000;
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                                                                                                                                                                            Sequence 316 AA;
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                                                                                                                                                                                                                            represents a human
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)B; AAS11687.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Page 39; 199pp; English.
                                                                          57;
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                                                                                                    Similarity
EARLHMPWYLLLGQLSLMDLLFTSVVTPKALADFLRRENTISFGGCALQMFLALTMG
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2000US-0180930P,
2000US-0181004P,
2000US-0181013P,
2000US-0181043P,
2000US-0220262P,
                                                                             Conservative
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2000US-0224596P.
2000US-0245292P.
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2000US-0180630P.
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                                                                                                    Score 57;
Pred. No.
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. 9.3e-44;
thes 0;
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                                                                                                                               Length 316;
                                                                                  Indels
                                                                                  0;
                                                                                  Gaps
                                  108
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Human GPCR polypeptide

SEQ ID NO 514

1

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AAU95574
ID AAU9
XX AAU9
AC AAU9
XX D2-2
XX D2-2
XX DE Huma
XX Huma
KW tran
KW ano1
KW schil
KW schil
KW schil
KW schil
KW aens
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KW schil
KW aegg1
XX Aug91
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Best Local S
Matches 57
                                                                 Human; olfactory and pheromone G protein coupled; receptor; GPCR; tranquilizer; antidepressant; neuroleptic; endocrine; anabolic; anorectic; taste; fragrance; food additive; cosmetic; cell migration; sterility; psychotic disorder; enurological disorder; anxiety; schizophrenia; manic depression; depression; axonal growth; menstrual cycle; appetite sexual motivation; sexual attraction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a method for screening G protein-coupled receptor (GPCR) genes (ABZ42870-ABZ43216) and/or GPCR proteins (ABP95596-ABP95942) by extracting open-reading frames containing 6-8 transmembrane domains with 250-1000 amino acid residues to give a gene homologous with studying in vivo signal transduction mechanism and identifying targets for drug development e.g. based on olfactory and gustatory receptors in form of agonists and antagonists by screening intrinsic and extrinsic improvers. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                           aggression
                                                                                                                                                                                                                                                                                          Human olfactory and pheromone G
                                                                                                                                                                                                                                                                                                                                                                  02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAU95574 standard;
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13-FEB-2001; 2001JP-00034434.
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evelopment; gustatory; taste; fragrance; receptor.
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Pred. No.
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Homo sapiens

Homo sapiens

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RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                      Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                        improvement, elimination or substitution of an existing taste and/or a fragrance of (or in) the food and/or cosmetic products. They can also be used in the preparation of medicament in the treatment and/or prevention of a mammalian disorder, such as cell migration, sterility, psychotic and depression, depression, for promoting anxiety, schizophrenia, manic connection and nerve regeneration for modulating male and female endocrine functions, hormone production and the menstrual cycle, for the behaviours, such as stimulation or suppression of appetite, sexual motivation, sexual attraction, aggression and for promoting or suppressing chemical communication between organisms. The present sequence is a human olfactory and pheromone GPCR protein sequence
                                          Human; olfactory G-coupled receptor; odour composition; taste composition.
                                                                                             G-coupled olfactory receptor #95.
                                                                                                                                    08-MAY-2002
                                                                                                                                                                      AAU85234;
                                                                                                                                                                                                      AAU85234 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    receptor (GPCR) or a protein 95% identical to the GPCR, a specific active portion and its encoding polymucleotide. Also included are an agonist, antagonist or inhibitor of the GPCR or the polymucleotide, a vector comprising the polymucleotide, a cell transformed by the vector, a non-human mammal comprising a partial or total deletion of the polymucleotide encoding the receptor and screening (detection and possibly, recovering) of compounds which are known or not known to be agonist, antagonists or inhibitors of natural compounds to the GPCR. The receptor-derived agonists, antagonists, inhibitors or compounds are used as an agonists, inhibitors or compounds are used as an
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 200
N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 197-198; 833pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel pheromone G-protein coupled receptor and receptor-derived agonists, antagonists or inhibitors useful in food or cosmetic products or in the treatment or prevention of neurological disorders such as anxiety and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Veithen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-SEP-2000; 2000EP-00870211.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-SEP-2001; 2001WO-BE000162.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention relates to olfactory and Pheromone G-protein coupled eptor (GPCR) or a protein 95% identical to the GPCR, a specific
                                                                                                                                                                                                                                                                                                       52 EARLHMEMYLLLGQLSLMDLLFTSVVTFKALADFLRRENTISFGGCALQMFLALTMG 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2002-330013/36.
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                                                                                                                                                                                                                                                                                                                                                                   57;
                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                     EARLHMPMYLLLGQLSLMDLLFTSVVTPKALADFLRRENTISFGGCALQMFLALTWG
                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                (first entry)
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                                                                                                                                                                                                      protein; 316 AA
                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                   18.0%;
                                                                                                                                                                                                                                                                                                                                                        Score 57; DB;
; Pred. No. 9.3;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                              0;
                                                     sensory perception of odourant;
                                                                                                                                                                                                                                                                                                                                                                           DB 5; Le
9.3e-44;
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                                                                                                                                                                                                                                                                                                                                                                                           Length 316
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RESULT 10
ADC85823
ID ADC85
XX ADC85
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XX ADC85
XX O1-JA
DT O1-JA
DX Humar
XX humar
XX Humar
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of representing sensory perception of one or more odourants. The method comprises: (a) providing a representative class of n olfactory receptors or ligand binding domains (LBD) of these receptors; (b) measuring values X1 to Xn representative of at least one activity of one or more odourants selected from: (i) binding one or more odourants to the LBD of at least one of the n olfactory receptors; (ii) activating at least one of the n olfactory receptors with the one or more odourants; and (iii) blocking at least one of the n olfactory receptors with the one or more odourants; and (c) generating a representation of sensory perception from the values X1 to Xn. The representation of the sensory perception of odourants is useful for the design and formulation of odour and taste compositions. ANU85140-ANU85393 designs the term of the values X1 to Xn. The representation of first of odourants is useful for the design and formulation of odour and taste compositions. ANU85140-ANU85393 designs the values X1 to Xn. The representation of odour and taste compositions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-JUN-2000; 2000US-0213812P.
13-MAR-2001; 2001US-00804291.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 101; 182pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 316 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                       human; GPCR; guanosine triphosphate-binding protein coupled receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human GPCR protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADC85823 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC85823;
                                                                                                                18-JUN-2001; 2001JP-00246789
                                                                                                                                                                                                                                          02-JAN-2003.
                                                                                                                                                                                                                                                                                                    EP1270724-A2
                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                             18-JUN-2002; 2002EP-00013517
                           (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY. (ADSC-) CENT ADVANCED SCI & TECHNOLOGY INCUBATIO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2002-083330/11.
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57; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein; 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO:276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Score 57; DB
%; Pred. No. 9.3
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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RESULT 11
AAUU7088
ID AAUU7088
AC AAUU7
XX AAUU7
AC AAUU7
XX AUU7
XX Humas
XX Humas
XX Humas
XX Humas
XX Hooct:
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WW mult
XX WO20
XX W020
XX O7-F
PR 07-F
PR 08-F
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Best Local
                                                                  07-FEB-2000; 2000US-0180646F.
08-FEB-2000; 2000US-0180930F.
08-FEB-2000; 2000US-0181004F.
08-FEB-2000; 2000US-0181013F.
08-FEB-2000; 2000US-0181013F.
08-FEB-2000; 2000US-0181043F.
24-UUL-2000; 2000US-0220262F.
25-UUL-2000; 2000US-0220594F.
11-AUG-2000; 2000US-0224596F.
02-NOV-2000; 2000US-0224596F.
06-FEB-2001; 2001US-00777789.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel polynucleotide encoding a guanosine triphosphate-binding protein coupled receptor (GPCR). A polynucleotide of the invention may have a use in gene therapy. The polynucleotide and polypeptide are useful for preparing a composition for treating a patient in need of increased or suppressed activity or expression of the guanosine triphosphate-binding protein coupled receptor. The protein sequences shown in ADC85549-ADC87617 represent GPCR's of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New polynucleotide, useful for preparing a composition for treating patient in need of increased or suppressed activity or expression guanosine triphosphate-binding protein coupled receptor.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; olfactory receptor; OR; G-protein coupled receptor superfamily; GPCR; gene therapy; tranquiliser; vulnerary; anti-HIV; cytostatic; nootropic; neuroprotective; antiparkinsonian; antiasthmatic; cancer; antiinflammatory; odorant receptor; olfactory loss; trauma; HIV; human immunodeficiency virus; neoplastic growth; neurological disorder; parkinson's disease; Alzheimer's disease; neurogenesis; Crohn's disease; wound healing; asthma; Albright hereditary osteodystrophy; multiple sclerosis; nasal epithelial neuronal tissue; transgenic plant.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 316 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human odorant receptor (OR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAU07088;
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                                                                                                                                                                                                                                                                                                    07-FEB-2000; 2000US-0180511P.
07-FEB-2000; 2000US-0180630P.
07-FEB-2000; 2000US-0180646P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200157215-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
                          (CURA-)
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57; Conserv
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                             CURAGEN
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100.0%; Pr
170 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ξ
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of the
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RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
WPI; 2002-471499/50
                                         Alsobrook JP,
                                                                                                                                                                                                                                                                         NOV; odorant receptor; G protein coupled receptor; GPCR; trauma; olfactory receptor; olfactory loss; neoplastic growth; human immunodeficiency virus; Alzheimer's disease; HIV; neurological disorder; Parkinson's disease.
                                                                                                02-NOV-2000; 2000US-0245292P
06-FEB-2001; 2001US-00777789
20-SEP-2001; 2001US-00245292
                                                                                                                                                         09-OCT-2001; 2001WO-US031744.
                                                                                                                                                                                           10-MAY-2002
                                                                                                                                                                                                                       WO200236632-A2
                                                                                                                                                                                                                                                     Unidentified
                                                                          (CURA-)
                                                                                                                                                                                                                                                                                                                                                       Novel odourant receptor NOV7
                                                                                                                                                                                                                                                                                                                                                                                       15-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU97928 standard; protein; 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel isolated NOVX polypeptide related to human odorant receptor family of G-protein coupled receptor superfamily of proteins useful for treating trauma, neoplastic growth, Parkinson's disease, Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Padigaru M,
Baumgartner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001-488883/53.
DB; AAS11688.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Page 44; 199pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   L Similarity
57; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BARLHMEMYLLLGOLSLMDLLFTSVVTPKALADFLRRENTISFGGCALQMFLALTMG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EARLHMPMYLLLGQLSLMDLLFTSVVTDKALADFLRRENTISFGGCALQMFLALTMG 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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JC;
                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ĀΑ;
                                         Burgess CE,
                                                                         CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18.0%;
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                                      Grosse
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                                                                                                                                                                                                                                                                                                                                                       protein.
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                                        ξ¥
                                     Lepley DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 4; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 324;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mishra VS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                     Padigaru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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RESULT 13
AAU97927
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC This invention relates to the DNA and protein sequences of a novel CC isolated olfactory receptor-like polypeptide, NOVX. The DNA and protein CC sequences of the invention and an antibody specific for the protein are CC useful for treating or preventing a disorder associated with NOVI in a CC subject, preferably human. A NOVX specific antibody is useful for CC protein and antibody of the invention is useful for diagnosing.

CC protein and antibody of the invention is useful for diagnosing.

CC protein and antibody of the invention is useful for diagnosing.

CC protein and antibody of the invention is useful for diagnosing.

CC protein and antibody of the invention is useful for diagnosing.

CC protein and solisorders associated with aberrant NOVX expression conclude the conclusion of the protein and an immunodeficiency parkinson's disease and Alzheimer's disease. The sequences and NOV contromosomal mapping, tissue typing, forensic biology), predictive contromosomal mapping, tissue typing, forensic biology), predictive conclusions (e.g., diagnostic assays, prognostic assays, monitoring clinical characterian of the protein and appropriate and prophylactic). The NOV protein is useful as immunogen to protential agonist and antagonist for the protein, to screen for the protein or three-hybrid assay. The nucleotide sequence is useful in gene in a NOVX gene, and to modulate NOVX activity. The antibody of the protein in useful for isolating, and purifying the protein and to consist or protein levels in tissue as part of a clinical testing procedure.

CC monitor protein levels in tissue as part of a clinical testing procedure.

The present sequence represents the NOV7 protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 57
                            02-NOV-2000; 2000US-0245292P.
06-FEB-2001; 2001US-00777789.
20-SEP-2001; 2001US-00245292.
                                                                                                                                                                                                                                                                           NOV; odorant receptor; G protein coupled receptor; GPCR; olfactory receptor; olfactory loss; neoplastic growth; human immunodeficiency virus; Alzheimer's disease; HIV;
                                                                                                             09-OCT-2001;
                                                                                                                                                    10-MAY-2002.
                                                                                                                                                                                                                               Unidentified.
                                                                                                                                                                                      WO200236632-A2
                                                                                                                                                                                                                                                                   neurological
                                                                                                                                                                                                                                                                                                                                                                  Novel
                                                                                                                                                                                                                                                                                                                                                                                                       15-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAU97927 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 324 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated olfactory receptor-like polypeptide, NOVX, useful for diagnosing, preventing or treating disorders of olfactory loss, e.g. trauma, human immunodeficiency virus illness, neoplastic growth and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 34-35; 122pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neurological disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             52 EARLHMEMYLLLGQLSLMDLLETSVVTPKALADFLRRENTISFGGCALQMELALTMG 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BARLHMPMYLLLGÓLSLMDLLFTSVVTPKALADFLRRENTISFGGCALÓMFLALTMG 116
                                                                                                         2001WO-US031744
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                               disorder; Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                          receptor NOV6 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18.0%; Score 57;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                324 AA
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hes 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5; Length 324;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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CURAGEN

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cc isolated olfactory receptor-like polypeptide, NOVX. The DNA and protein cc isolated olfactory receptor-like polypeptide, NOVX. The DNA and protein cc sequences of the invention and an antibody specific for the protein are useful for treating or preventing a disorder associated with NOVI in a customer of protein and antibody is useful for cc gubject, preferably human. A NOVX specific antibody is useful for cc preventing or treating disorders associated with NOVI in a cc preventing or treating disorders associated with NOVI in a sample. The DNA, cc preventing or treating disorders associated with aberrant NOVX expression cc preventing or treating disorders e.g. trauma, human immunodeficiency cr wirus (HIV) illness, neoplastic growth and neurological disorders, e.g. cr ratibody is useful in screening assays, detection assays (e.g., antibody is useful in screening assays, detection assays, predictive cc medicine (e.g., diagnostic assays, prognostic assays, monitoring clinical cr rials and pharmacogenomic), and in methods of treatment (e.g., cc produce antibodies immunospecific for the protein is useful as immunospec cc protein and antagonist compounds, and as bait protein in a two-thorid or three-hybrid assay. The nucleotide sequence is useful in gene therapy, to express the protein, to detect NOVX mRNA or a genetic lesion in a NOVX gene, and to modulate NOVX activity. The antibody of the compounds and antagonist compounds and antagonist compounds and antagonist compounds. The protein in a modulate note in a section of the invention of the protein levels in tissue as part of a clinical testing procedure.

Cc invention is useful for isolating, and purifying the protein and to compound to compound to the protein levels in tissue as part of a clinical testing procedure.

Cc invention sequence represents the NOV6 protein of the invention
ARBSULT 14
AAG773
ID AAG773
XX BECOTO
XX BECOTO
XX Unida
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 33; 122pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neurological disorders.
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                                                                                                                                                                                                                                                                                                                                    Human; olfactory receptor; OR; primary scent determination; secondary scent determination; polypeptide library; odour receptor; scent profile; scent fingerprint; scent representation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Olfactory receptor-like polypeptide, SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAG73027 standard;
                                                                                                                                                                                                                                                                      Unidentified
      08-OCT-1999;
                                                                        06-OCT-2000;
                                                                                                                                      19-APR-2001.
                                                                                                                                                                                                        WO200127158-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2002-471499/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK53096.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                           2000WO-US027582.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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          99US-0158615P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18.0%; Score 57;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Grosse WM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      234 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 5; Le
9.5e-44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 324;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2709.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          y loss, e.g.
growth and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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RESULT 15
AAB71196
ID AAB71
XX AAB71
XX T2-NC
CX AAB71
XX Humar
XX 101-D
XX 118-D
PR 21-D
PR 21-D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is an olfactory receptor polypeptide which was used CC as a query sequence in a database search of olfactory receptor (OR)-like CC sequences. The invention relates to isolated polynucleotides encoding CC polypeptides involved in olfactory sensation. The polynucleotides can be CC used in screening for olfactory agonists and antagonists. The methods CC used in screening for olfactory agonists and antagonists. The methods of CC combination of primary scents and the identification of CC enable determination of secondary scents and the identification of CC combinations of odour receptors used to detect these primary scents. The methods also CC combinations of odour receptors that are involved in detecting such CC secondary scents. This enables the construction of a scent representation CC (also called a scent fingerprint or scent profile), which may be used to CC (also called scents. Libraries of olfactory receptors are useful CC receptors, and can be used for determining differences in the olfactory receptors are useful constructions of a scent receptors are useful constructions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-290713/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bellenson J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-FEB-2000; 2000US-0184809P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 6; Page 1832-1833; 1857pp; English.
                                                                                                                                                                                                                                                         Human; anti-HIV; nootropic; antiasthmatic; antiarteriosclerotic; GPCRX; immunosuppressive; immunomodulator; cytostatic; antiinflammatory; AIDS; antidiabetic; neuroprotective; anorectic; haemostatic; antibacterial; fungicide; protozoal; virucide; human G-protein coupled receptor; gene therapy; vaccine; cardiomyopathy; atherosclerosis; diabetes; cell signal processing; cancer; obesity; neurodegenerative disorder; cachexia; anorexia; Alzheimer's disease; parkinson's disease; immune disorder; graft versus host disease; bronchial asthma; crobn's disease; multiple sclerosis; haemophilia; infectious disease; infectious disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 234 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human GPCRX protein SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB71196 standard; protein; 316 AA.
                                                                                                                                                                                                       Homo sapiens.
                                                                                 18-DEC-2001; 2001WO-US048958.
                                                                                                                         27-JUN-2002.
                                                                                                                                                                                                                                            diopathic thrombocytopenic purpura; receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polynucleotides which encode polypeptides sation for identifying olfactory agonists a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 118
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YEDA RES & DEV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        68.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 4; Le
2.7e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and antagonists
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      involved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fuchs T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 234;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          in olfactory
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18-DEC-2000; 21-DEC-2000; 04-JAN-2001;

2000US-0256635P. 2000US-0257876P. 2001US-0259743P.

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                                                                       Query Match
Best Local S
Matches 15
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12-JAN-2001; 2001US-0261498P

24-JAN-2001; 2001US-0267464P

08-FEB-2001; 2001US-0267464P

22-FEB-2001; 2001US-0277021P

14-MAR-2001; 2001US-0279946P

23-MAR-2001; 2001US-02798150P

18-JAPR-2001; 2001US-0284591P

23-APR-2001; 2001US-0285718P

19-JUN-2001; 2001US-039327P

16-AUG-2001; 2001US-0312902P
                                                                Sequence 316 AA;
                                                                                                                                                                                                                                                                                    This invention describes novel human G-protein coupled receptor (GPCRX) polypeptides which have anti-HIV, nootropic, antiasthmatic, protozoal, antiarteriosclerotic; immunosuppressive, immunomodulator, cytostatic,
                                                                                                                                                                                                                                                                                                                                    New G-protein coupled receptor polypeptides, useful in gene therapy, particularly for treating or preventing cardiomyopathy, atheroscierosis, diabetes, Crohn's disease, hemophilia or cancer in humans.
                                                                                                                                                                                                                                                                                                                       Claim 1; Page 112; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                 Padigaru M, Kekuda R,
Baumgartner JC, Burges
                                                                                                                                                                                                                                                                                                                                                                                                                           (CURA-) CURAGEN CORP.
124 MAYDRYVAICHPLKY 138
               118 МАУДКУVАІСНРІКУ 132
                                15;
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                                        Similarity
                                Conservative
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ss CE;
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                             Score 15; DB 5; Lo
Pred. No. 3.5e-05;
0; Mismatches 0;
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Search completed: October Job time: 127 secs

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316
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Copyright (c) 1993 - 2004 Compugen Ltd.
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ALIGNMENTS

10-Nov-1995 #text_change 04-Sep-1998

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A;Molecule type: mRNA
A;Residues: 1-232 <ABE>
C;Superfamily: olfactory receptor OR14
C;Keywords: G protein-coupled receptor; transmembrane protein
                                                                                                                                                                                                          R;Abe, K.; Kusakabe, Y.; Tanemura, K.; Emori, Y.; Arai, S. FEBS Lett. 316, 253-256, 193 A;Title: Multiple genes for G protein-coupled receptors and their expression in lingual A;Reference number: S28996; MUID:93138133; PMID:8380780 A;Accession: S29001
                                                                                                                                                                                                                                                                                                                                                                                          RESULT 2
S29001
                                                                                                                                                                                                                                                                                             R;Abe,
                                                                                                                                                                                                                                                                                                   G protein-coupled receptor (clone PTE58) - rat (fragment) C;Species: Rattus norvegicus (Norway rat) C;Date: 25-Feb-1994 #sequence_revision 10-Nov-1995 #text_change C;Accession: S29001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1

$28999

G protein-coupled receptor (clone PTE38) - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_(C;Accession: $28999
R;Abe, K, Kusakabe, Y:; Tanemura, K.; Emori, Y.; Arai, S
FEBS Lett. 316, 253-256, 1993
A;Title: Multiple genes for G protein-coupled receptors a:
A;Reference number: $28996; MUID:93138133; PMID:8380780
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Superfamily: olfactory receptor OR14
C;Keywords: G protein-coupled receptor; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-234 < ABE>
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                                                                                      Query Match
Best Local Similarity
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                          118 MAYDRYVAICHPL 130
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58 MAYDRYVAICHPL 70
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Pred. No. 9.1e-
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                                                                  Mismatches
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R;Thomas, M.B.; Haines, S.L.; Akeson, R.A.
Gene 178, 1-5, 1996
A;Title: Chemoreceptors expressed in taste,
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C;Date: 27-Feb-1997 #sequence revision 27-Feb-1997 #text_change 26-Aug-1999
C;Accession: JC5200; PC4302
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A;Accession: PC4302
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A; Accession: JC5200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Gene: tb334 (C;Superfamily: Olfactory receptor OR14 (C;Superfamily: Olfaction; taste bud; transmembrane protein C;Keywords: olfaction; taste bud; transmembrane predicted <TM1>F;24-4-47)Domain: transmembrane #status predicted <TM2>F;56-77/Domain: transmembrane #status predicted <TM2>F;56-77/Doma
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F;138-162/Domain: transmembrane #status predicted <TM4>
F;195-217/Domain: transmembrane #status predicted <TM5>
F;236-258/Domain: transmembrane #status predicted <TM6>
F;236-258/Domain: transmembrane #status predicted <TM6>
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A; Residues: 146-153; 265-272 < THO2>
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A;Status: preliminary
A;Molecule type: protein
A;Rolecule type: protein
A;Rolecule type: protein
A;Rosidues: 144-151;270-277 <THO2>
A;Rosidues: 144-151;270-277 <THO2>
A;Rosidues: 144-151;270-277 <THO2>
A;Rosperimental source: taste bud
C;Comment: This protein is coupled to a GTP-binding protein-mediated signal transduction
C;Genetics:
A;Genetics:
A;Genetics:
A;Genetics:
C;Genetics:
A;Genetics:
A;Gen
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JC5201
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C;Date: 20-Feb-1997 #sequence_revision 27-Feb-1997 #text_change 26-Aug-1999
C;Accession: JC5201; PC4303
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A;Accession: PC4303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Status: preliminary; nucleic acid sequence not shown A;Molecule type: DNA A;Residues: 1-315 <THO1>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Title: Chemoreceptors expressed in taste, A;Reference number: JC5200; MUID:97080538; lA;Accession: JC5201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Thomas, M.B.; Haines, S.L.; Akeson, R.A. Gene 178, 1-5, 1996
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nilarity 100.0%;
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Pred. No.
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R;Buck, L.; Axel, R.

Cell 65, 175-187, 1991

A;Title: A novel multigene family may encode odorant receptors:
A;Reference number: A23701; MUID:91191556; PMID:1840504

A;Accession: F23701

A;Accession: F23701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  olfactory receptor I7 - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change 26-Aug-1999
C;Accession: F23701
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F23701
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A;Residues: 1-327 <BUC>
A;Cross references: GB:M64386; NID:g205833; PIDN:AAA41749.1; PID:g205834
C;Superfamily: olfactory receptor OR14
C;Keywords: G protein-coupled receptor; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Status: nucleic acid sequence not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Buck, L.; Axel, R. Cell 65, 175-187, 1991
Cell 65, 175-187, 1991
A;Title: A novel multigene family may encode odorant receptors: a A;Reference number: A23701; MUID:91191556; PMID:1840504
A;Accession: A23701
A;Accession: A23701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 olfactory receptor F3 - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change 26-Aug-1999
C;Accession: A23701
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A;Residues: 1-333 <BUC>
A;Residues: 1-333 <BUC>
A;Cross-references: GB:M64376; NID:g205813; PIDN:AAAA41739.1; PID:g205814
C;Keywords: G protein-coupled receptor; transmembrane protein
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odorant receptor (clone K4) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 22-Sep-1993 #sequence_revision 18-Nov-1994 #text_change 03-No
C;Accession: D40745
R;ResBler, K.J.; Sullivan, S.L.; Buck, L.B.
Cell 73, 597-609, 1993
A;Title: A zonal organization of odorant receptor gene expression in
A;Reference number: A40745; MUID:93258822; pMID:7683976
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Status: nucleic acid sequence not shown
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100.0%; Pred. No.
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                                                                                                         #text_change 03-Nov-2000
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A;Cross-____C;Superfamily:
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A; Residues: 1-309 < GAT>
A; Residues: 1-309 < GAT>
A; Cross-references: EMBL: X80671; NID: g517365; PIDN: CAA56697.1; PID: g517366
R; Gat, U.; Nekrasova, E.; Lancet, D.; Natochin, M.
submitted to the EMBL Data Library, July 1994
submitted to the EMBL Data Library, July 1994
                                                                                                                                   A; Molecule type: mRNA
A; Residues: 1-309 <GA2>
                                                                                                                                                                                                                                                                                                                                                                            R;Gat, U.; Nekrasova, E.; Lancet, D.; Natochin, Eur. J. Biochem. 225, 1157-1168, 1994
A;Title: Olfactory receptor proteins. Expression
                                                                                                                                                                           A;Status: preliminary
                                                                                                                                                                                                      A; Description: Olfactory receptor proteins: A; Reference number: $47014
                                                                                                                                                                                                                                                                                                                                                 A; Reference number: S51356; A; Accession: S51356
                                                                                                                                                                                                                                                                                                                                                                                                                        C;Species: Rattus norvegicus (Norway rat)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession: S51356; S47014
                                                                                                                                                                                             A; Accession: S47014
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Title: Cloning and expression of odorant A; Reference number: S29707; MUID:93149273; A; Accession: S29709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Date: 25-Peb-1994 #sequence_revision 10-Nov-1995 #text_change 04-Sep-1998 C;Accession: S29709
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A;Exter sequence extracted from NCBI backbone (NCBIP:131753)
C;Superfamily: olfactory receptor OR14
C;Keywords: G protein-coupled receptor; transmembrane protein
                                                   Query Match
Best Local Similarity
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A;Status: preliminary; not compared with conceptual translation A;Molecule type: nucleic acid
A;Residues: 1-222 <RES>
                                                                                           Cross-references: EMBL:X80671; NID:g517365; PIDN:CAA56697.1; PID:g517366 Superfamily: olfactory receptor OR14
                                                                                                                                                                                                                                                                                                                                    Status: preliminary
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Best Local
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3.5%; oc.
100.0%; Pred. No.
-ive 0; Mismatches
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MUID:95045546; PMID:7957207
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0.011;
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PMID:7678922
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C;Accession: 138473
R;Ben-Arie, N.; Lancet, D.; Taylor, C.; Khen, M.; Walker, N.; Hum. Mol. Genet. 3, 229-235, 1994
A;Title: Olfactory receptor gene cluster on human chromosome 1
A;Reference number: 138470; MUID:94272458; PMID:8004088
A;Accession: 138473
A;Accession: 138473
                                                                                                                                                                                                                                              RESULT 12
I38473
                        A; Molecule type: DNA
A: Residues: 1-215 < RES>
                                                                                                                                                                                                    olfactory receptor - human (fragment) C;Species: Homo sapiens (man)
                                                                                                                                                                    C;Species: Homo sapiens (man)
C;Date: 29-May-1998 #sequence_revision 29-May-1998 #text_change 26-Aug-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Superfamily: olfactory receptor OR14
C;Keywords: G protein-coupled receptor; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Abe, K.; Kusakabe, Y.; Tanemura, K.; Emori, FEBS Lett. 316, 253-256, 1993
A;Title: Multiple genes for G protein-coupled A;Reference number: 828996; MUID:93138133; PMI A;Accession: $28997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: mRNA
A; Residues: 1-168 < ABE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Date: 25-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 04-Sep-1998 C;Accession: S28997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 11
S28997
;Cross-references: EMBL:U04681; NID:g438395; PIDN:AAA18345.1; PID:g438396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              G protein-coupled receptor (clone PTEO3) - rat
C;Species: Rattus norvegicus (Norway rat)
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R;Ben-Arie, N.; Lancet, D.; Taylor, C.; Khen, M.; Walker, Hum. Mol. Genet. 3, 229-235, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Gene: OR17-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Title: Olfactory receptor gene cluster on human chromosome 17: possible duplication A;Reference number: I38470; MUID:94272458; PMID:8004088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Residues: 1-165 <RES>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
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C;Genetics:
A;Gene: OR17-30
C;Superfamily: olfactory receptor OR14
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C; Superfamily:
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R;Ben-Arie, N.; Lancet, D.; Taylor, C.; Khen, M.; Walker, N.; Ledbetter, D.H.; Carrozzo, Hum. Mol. Genet. 3, 229-235, 1994
A;Title: Olfactory receptor gene cluster on human chromosome 17: possible duplication of A;Reference number: I38470; MUID:94272458; PMID:8004088
A;Accession: I38470
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C;Species: Homo sapiens (man)
C;Date: 29-May-1998 #sequence_revision 29-May-1998 #text_change 26-Aug-1999
C;Accession: I38470
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A; Residues: 1-216 < RES>
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R;Ben-Arie, N.; Lancet, D.; Taylor, C.; Khen, M.; Walker, N.; Ledbetter, D.H.; Carrozzo, Hum. Mol. Genet. 3, 229-235, 1994
A;Title: Olfactory receptor gene cluster on human chromosome 17: possible duplication of A;Reference number: I38470; MUID:94272458; PMID:8004088
A;Accession: I38474
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A;Molecule type: DNA
A;Residues: 1-216 <RES>
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                                        RESULT 15
C40745
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A;Gene: OR17-32
C;Superfamily: olfactory receptor OR14
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C;Genetics:
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Best Local Similarity
odorant receptor (clone K7) - mouse (fragment) C;Species: Mus musculus (house mouse)
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tive 0; Mismatches
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C;Date: 22-Sep-1993 #sequence_revision 18-Nov-1994 #text_change 26-Aug-1999 C;Accession: C40745 R;Ressler, K.J.; Sullivan, S.L.; Buck, L.B.

Cell 73, 597-609, 1993 A;Title: A zonal organization of odorant receptor gene expression in the olf A;Reference number: A40745; MUID:93258822; PMID:7683976 A;Accession: C40745 A;Accessi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: GB:L14569; NID:g293759; PIDN:AAA39853.1; PID:g293760
A;Experimental source: olfactory epithelium
A;Note: sequence extracted from NCBI backbone (NCBIP:131750)
C;Superfamily: olfactory receptor OR14
C;Keywords: G protein-coupled receptor; transmembrane protein
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Similarity 100.0%; Pred. No.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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man Oli	IGS to :	. U.S.A. /e odoran	orthol	r., You	ROM N.	2002).	scheme	R., St	Malecova P	to the		ni H.,	ouchi I.		Chordata; Primates;	•		, Created)	ARD;			315 1 322 1		311 1					324 L 327 1	
DATABASE: NAME=Human Olfactory Receptor Data Exploratorium (HORDE); WWW="http://bioinformatics.weizmann.ac.il/cgi-bin/HORDE/showGene.nl?kev=sv	SUBCELLULAR LOCATION: Integral SIMILARITY: Belongs to family 1	U.S.A. 98:7390-73 odorant receptor.	nalysis of orthologous mouse and l		A. 416212;		for deci	다;		EMBL/GenBank/DDBJ		Asai K.,	., Arita				Last sequence uplast annotation (HT3).	ed)	PRT;	ALI		05A1_H 05P2_H	OR23 M	05AN_H	OAV1_H	05K3_H	07G1_H	O6B3_HUMAN	O7.F7_R	
Recepto mann.ac	membrane L of G-pr	98:7390-7395(2001) receptor.	use and	Athanasiou M.			deciphering	M., O'Brien	2	:nBank/E		Akiyama Y.;	. м., Fo		Craniata; Ve Catarrhini;		e update) lon update)		316 /	ALIGNMENTS		HUMAN	MOUSE	HUMAN	HUMAN	HUMAN	UMAN	UMAN	HUMAN	
r Data .il/cqi	e protein. rotein cou	(2001).	numan				families	rien J.K.,	3			la Y.;	ıtami K.,		Vertebrata; E i; Hominidae;		e) ate)		AA.											
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orium	recept			n C.,			genes.";	Radelof U.,			T attamenent atte		co.		Euteleostomi; Homo.							Q8ngjo 1 Q8wz92 1							Q8ng99 h	
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WWW-"http://bioinformatics.weizmann.ac.il/cgi-bin/HoRDE/showdene.pl?key=symbols.

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CC or send an email to license@isb-sib.ch).

CC EMBL; AB065823; BAC06042.1; ...

DR EMBL; AF39518; AAWS5103.1; ...

DR EMBL; AF3912137; AAG45209.1; ...

DR EMBL; AF321237; AAG45209.1; ...

DR Genew; HGNC:15142; ORZAGI.

DR Ffam; PF00001; 7tm_1; 1.

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01-JUN-1994 (Rel. 29, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
30-MAY-2000 (Rel. 39, Last concepts)
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P35897;
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Rattus norvegicus (Rat)
Eukaryota, Metazoa, Chordata, Cran
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CARBOHYD
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STRAIN=Fischer; TISSUE=Lingual epithelium;
MEDLINE=9313813; PubMed=8380780;
MEDLINE=9313813; PubMed=8380780;
Abe K., Kusakabe Y., Tanemura K., Emori Y., Arai
"Multiple genes for G protein-coupled receptors"
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PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus
                                                                                                                               G-protein
NON_TER
                                                                                                                                                                                                    -!- FUNCTION: Possible taste receptor.
-!- SUBCELLULAR LOCATION: Integral membrane protein.
-!- TISSUE SPECIFICITY: Tongue specific.
-!- SIMILARITY: Belongs to family 1 of G-protein coupled receptors.
InterPro, IPR000276, GPCR Rhodpsn.
Pfam; PF00001; 7tm 1; 1.
                                                                                                                                                                                                                                                                                                           in lingual epithelia.";
FEBS Lett. 316:253-256(1993).
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57; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EARLHMPMYLLLGQLSLMDLLFTSVVTPKALADFLRRENTISFGGCALQMFLALTMG 108
                                                                                                                                                                  PS00237; G PROTEIN_RECEP_F1_1; 1.
PS50262; G_PROTEIN_RECEP_F1_2; 1.
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GPCRRHODOPSN,
GPROTEIN RECEP_F1_2; 1.
GPROTEIN_RECEP_F1_2; 1.
dreceptor; Transmembrane; Glycoprotein;
Olfaction.
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N-LINKED (GLCNAC. . .)
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                                                                                                                                                                                                                                                                                                                                                                                                    receptor genes.";
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                           Suwa M., Sato T., Okouchi I., Arita M., Futami K., Matsumoto S.,
Tsutsumi S., Aburatani H., Asai K., Akiyama Y.;
"Genome-wide discovery and analysis of human seven transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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                                                                                                                                                                    PRINTS; PR00237; GFCRRHODOPSN.
PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; 1.
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
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Pfam; PF00001; 7tm_1; 1.
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SIMILARITY: Belongs to family 1 of G-protein coupled receptors.
DATABASE: NAME=Human Olfactory Receptor Data Exploratorium (HORDE);
WWW="http://bioinformatics.weizmann.ac.il/cgi-bin/HORDE/showGene.pl?key=symbol&v
                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION: Putative odorant receptor.
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60 MAYDRYVAICHPLKY 74
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Pred. No.
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EXTRACELLULAR (POTENTIAL).

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                                                                                                                                                                                                                                                                        noved. Usage by and for commercial (See http://www.isb-sib.ch/announce/
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4.5e-07;
                                                                                                                                                             Glycoprotein;
                                                 (POTENTIAL)
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RESULT 4
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Olfaction.
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InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the European Bioinformatics Institute. The use by non-profit institutions as long a modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                          Pfam; PF00001; 7tm 1; 1.

PROSITE; PS00237; G PROTEIN RECEP F1 1; 1.

PROSITE; PS50262; G PROTEIN RECEP F1 2; 1.
                                                                                                                                                                                                                                                                                                                                                                                        EMBL; U86216; AAC39612.1; -. Genew; HGNC:8221; OR1N1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rouquier S., Taviaux S., Trask B.J., Brand Demaille J.G., Giorgi D.; "Distribution of olfactory receptor genes"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=98160182; PubMed=9500546;
MEDLINE=98160182; PubMed=9500546;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    O1N1 HUMAN
O43870;
                                                                                                                                                       DOMAIN
                                                                                                                                                                                                                               DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                              or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                               entities requires a license agreement
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-OCT-2003 (Rel. 42, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
Olfactory receptor 1NI (Olfactory receptor 1-
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10-OCT-2003 (Rel.
15-MAR-2004 (Rel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUBCELLULAR LOCATION: Integral membrane protein.
SIMILARITY: Belongs to family 1 of G-protein coupled receptors.
DATABASE: NAME=Human Olfactory Receptor Data Exploratorium (HORDE);
WWW="http://bioinformatics.weizmann.ac.il/cgi-bin/HORDE/showdene.pl?key=symbol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FUNCTION: Putative odorant receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genet. 18:243-250(1998)
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100.0%; Pr
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7 (POTENTIAL).
BY SIMILARITY.
                                                                                         5 (POTENTIAL)
CYTOPLASMIC ()
6 (POTENTIAL)
                                                                                                                                                                                                                                                                                          Transmembrane; Multigene family;
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3 (POTENTIAL).
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0; Mismatches
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CYTOPLASMIC (P
BY SIMILARITY.
N-LINKED (GLCN
E27DA9AE1E2C50BE CRC64
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72D856594467060F CRC64;
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5.6e-06;
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Best Local
                                                                        EMBL; AF399617; AAX95102.1; EMBL; X89666; CAA61813.1; -- EMBL; X64992; --; NOT_ANNOTAT
                                                                                                                  modified and this statement is not removed. entities requires a license agreement (See or send an email to license@isb-sib.ch).
                                                              PIR; S58011; S58011
                                                                                                                                                                                                     Parmentier M., Libert F., Schurmans S., Schiffmann S., Lefort A., Eggerickx D., Ledent C., Mollereau C., Gerard C., Perret J., Grootegoed A., Vassart G.; "Expression of members of the putative olfactory receptor gene family in mammalian germ cells.";
                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
                                                                                                                                                              the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                    Nature 355:453-455(1992).
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SEQUENCE FROM N.A.
MEDLINE=22202146; PubMed=12213199;
Puchs T., Malecova B., Linhart C., Shar
Puchs T., Malecova R., Steinfath M.,
                                                                                                                                                                                                                                                                                                                                    MEDLINE=92131132; PubMed=1370859;
                                                                                                                                                                                                                                                                                                                                                          SEQUENCE OF 59-172 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                              cells of
                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=97224452; PubMed=9119360;
                                                                                                                                                                                                                                                                                                                                                                                    Genomics
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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10-OCT-2003 (Re
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                                                                                                                                                                                                                                                                                                                                                                                 Anderhaeghen P., Schurmans S., Vassart G., 
'Specific repertoire of Olfactory receptor 
'ells of several mammalian species."; 
enomics 39:239-246(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lehrach H., Lancet D., DEFOG: a practical sc
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15-MAR-2004 (Rel. 43,
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                                                                                                                                                                                                                                                     FUNCTION: Putative odorant receptor.
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Last annotation update)
(Olfactory receptor TPCR100)
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4e-05;
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                                                                                                                                       Usage
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in J.K.,
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.K., Radelof U.,
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InterPro; IPR000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm 1; 1.
PRINTS; PR000237; GECRRHODDENN.
PROSITE; PS00237; G_PROTEIN_RECEP_F1_1;
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2;

Genew; HGNC:8270;

OR2M4

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Best Local 9
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P35899;
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01-JUN-1994 (Rel. 29, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation updat
Possible gustatory receptor clone PTE58 (Fr
Rattus norvegicus (Rat)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=9313813; Pubmed=8380780; Arai S.; Abe K., Kusakabe Y., Tanemura K., Emori Y., Arai S.; "Multiple genes for G protein-coupled receptors and their in lingual epithelia."; Free Tett 37.000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                      PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
  DISULFID NON TER SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PIR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                            NON
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InterPro; IPR000276; GPCR_Rhodpsn.
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                                                                                                                                                                                                                                                                                                                                            PS00237; G_PROTEIN_RECEP_F1_1;
PS00237; G_PROTEIN_RECEP_F1_2;
PS50262; G_PROTEIN_RECEP_F1_2;
PS50262; G_PROTEIN_RECEP_F1_2;
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                                                      STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EXTRACELLULAR (POTENTIAL) 5 (POTENTIAL).
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7 (POTENTIAL)
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EXTRACELLULAR (POTENTIAL)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CYTOPLASMIC (POTENTIAL) .
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EXTRACELLULAR (POTENTIAL)
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EXTRACELLULAR (POTENTIAL)
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            2B72EEAB08A992B7 CRC64;
                                                                                                                                                                                (POTENTIAL)
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3. 4e-05;
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                                                                                                                                                                                                                                                               G-protein Olfaction.
                                                                                                                                                                                                                                                                                                              PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                   Genew; HGNC:15109; ORIJ5.
InterPro; IPR000276; GPCR_Rhodpsn.
                                                                                                                                             DOMAIN
                                                                                                                                                                   TRANSMEM
                                                                                                                                                                                                                                                                                                                                                             PF00001;
                                                                                                                                                                                                                                                                           ps00237; G_PROTEIN_RECEP_F1_1; 1.
ps00237; G_PROTEIN_RECEP_F1_2; 1.
ps50262; G_PROTEIN_RECEP_F1_2; 1.
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1 (POTENTIAL).

CYTOPLASMIC (POTENTIAL).

2 (POTENTIAL).

EXTRACELLULAR (POTENTIAL).

3 (POTENTIAL).

CYTOPLASMIC (POTENTIAL).

4 (POTENTIAL).

Multigene

EXTRACELLULAR (POTENTIAL) 5 (POTENTIAL).

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE OF 91-252 FROM N.A.
MEDLINE=20183981; PubMed=10706615;
Rouquier S., Blancher A., Giorgi D.;
"The olfactory receptor gene repertor of the feature of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                              entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in mammalian germ cells.";
Nature 355:453-455(1992).
-i- FUNCTION: Putative odorant receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "The olfactory receptor gene repertoire in primates a Evidence for reduction of the functional fraction in Proc. Natl. Acad. Sci. U.S.A. 97:2870-2874(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Parmentier M., Libert F., Schurmans
Eggerickx D., Ledent C., Mollereau
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=92131132; PubMed=1370859;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Expression of members of the putative olfactory receptor gene family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE OF 93-206 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 100
les 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SUBCELLULAR LOCATION: Integral membrane protein.
SIMILARITY: Belongs to family 1 of G-protein coupled receptors.
DATABASE: NAME=Human Olfactory Receptor Data Exploratorium (HORDE);
WWW="http://bioinformatics.weizmann.ac.il/cgi-bin/HORDE/showGene.pl?key=symbol&v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               118 MAYDRYVAICHPL 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 58 MAYDRYVAICHPL 70
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(Rel. 42, Last sequence update)
(Rel. 43, Last annotation update)
ceptor 1J5 (HSA5) (HTPCRXI5) (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (Human)
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NOT_ANNOTATED_CDS
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Primates;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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                                                                                                                                                        G-protein
Multigene
                                                                                                                                                                                                                                  EMBL; AB065496; BAC05747.1; -.
                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
                                                                        DOMAIN
                                                                                                                                                                            InterPro; IPR000276; GPCR Rhodpsn.
Pfam; PF000001; 7tm 1; 1.
PROSITE; PS00237; G PROTEIN RECEP_F1_1;
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2;
                                                                                                                                                                                                                                                                                                                                                                                                           receptor genes.",
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                            Genew; HGNC:15251; OR5AK2.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Tsutsumi S., Aburatani H., "Genome-wide discovery and
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                           FUNCTION: Putative odorant receptor.
SUBCELLULAR LOCATION: Integral membrane protein.
SIMILARITY: Belongs to family 1 of G-protein coupled receptors.
DATABASE: NAME-Human Olfactory Receptor Data Exploratorium (HOF
                                                                                                                                                                                                                                                                                                                                                  WWW="http://bioinformatics.weizmann.ac.il/cgi-bin/HORDE/showGene.pl?key=symbol
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in coupled receptor; Transmembrane; G:
ne family; Olfaction.

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M 26 46 1 (POTENTIAL).
47 54 CYTOPLASMIC (POTENTIAL).

M 55 75 2 (POTENTIAL).

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M 134 154 4 (POTENTIAL).

M 135 196 EXTRACELLULAR (I
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M 197 217 STOPLASMIC (POTENTIAL).

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6 (POTENTIAL).

EXTRACELLULAR (POTENTIAL).

7 (POTENTIAL).

CYTOPLASMIC (POTENTIAL).

BY SIMILARITY.

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G-protein Multigene DOMAIN TRANSMEM

PRINTS; PRO0237; GPCRHODOPSN PROSITE; PS00237; G PROTEIN RI PROSITE; PS50262; G PROTEIN RI

P800237; G PROTEIN_RECEP_F1_1; 1.
P850262; G_PROTEIN_RECEP_F1_2; 1.
n coupled receptor_Transmembrane;
e family; Olfaction.

Glycoprotein;

DOMAIN TRANSMEM

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EXTRACELLULAR (POTENTIAL)

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CYTOPLASMIC (POTENTIAL).

2 (POTENTIAL).

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                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Ew the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-s
                                                                                                                                                                                                                              Genew; HGNC:8469; OR8A1
                                                                                                                                                                                                                                           EMBL; AB065833; BAC06052.1; EMBL; AF399512; AAK94997.1;
                                                                                                                                                                                                                                                                                                              entities requires a license agreement (Some send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -!- SUBCELLULAR LOCATION: Integral membrane protein.
-!- SIMILARITY: Belongs to family 1 of G-protein coupled receptors
-!- DATABASE: NAME=Human Olfactory Receptor Data Exploratorium (HO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shmulevich D., Elkon R., Steinfa
Lehrach H., Lancet D., Shamir R.
"DEFGG: a practical scheme for d
Genomics 80:295-302(2002).
                                                                                                                                                                                    InterPro; IPR000276; GPCR_Rhodpsn
Pfam; PF00001; 7tm_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            receptor genes.";
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ
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Suwa M., Sato T., Okouchi I., Arita
Suwa M., Sato T., Okouchi I., Arita
Tsutsumi S., Aburatani H., Asai K.,
"Genome-wide discovery and analysis
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Mammalia; Eutheria;
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CYTOPLASMIC (POTENTIAL).

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M., O'Brien
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                                                                                                                                                                                                                                                                                                                                  (See http://www.isb-sib.ch/announce/
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n J.K., Radelof U.,
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                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                  Fuchs T., Malecova B., Linhart C. Shmulevich D., Elkon R., Steinfat Lehrach H., Lancet D., Shamir R.,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (Human).
                                                                                                                                                                                                                                                                           "DEFOG: a practical scheme for deciphering families of genes.", Genomics 80:295\text{-}302\,(2002).
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Tsutsumi S., Aburatani H., Asai K., Akiyama Y.,
"Genome-wide discovery and analysis of human seven
                                                                                                                                                                                                                                                                                                                                                         SEQUENCE OF 68-282 FROM N.A.
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pfam; pf00001; 7tm_1; 1.
prints; pr00237; Gperrecoopsn.
prosits; ps00237; G_protein_recep_f1_1; 1.
                                     EMBL; AB065734; BAC05955.1; -.
EMBL; AF399532; AAK95017.1; -.
Genew; HGNC:14724; OR901.
InterPro; IPR000276; GPCR_Rhodpsn
                                                                                                        entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                            MEDLINE=22202146; PubMed=12213199;
                                                                                                                                                                                                                                                                                                                                                                                               receptor genes.";
                                                                                                                                                                                                            SUBCELLULAR LOCATION: Integral membrane protein.
SUBCELLULAR LOCATION: Integral membrane protein coupled receptors.
SIMILARITY: Belongs to family 1 of G-protein coupled receptors.
DATABASE: NAME=Human Olfactory Receptor Data Exploratorium (HORDE);
WWW="http://bioinformatics.weizmann.ac.il/cgi-bin/HORDE/showGene.pl?key=symbol
                                                                                                                                                                                                                                                                 FUNCTION: Putative odorant receptor.
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QBNGSO, Q96R93;
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15-MAR-2004 (Rel. 43, I
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                                                                                                                                                                                                       Fuchs T., Malecova B., Linhart C., Sharan R., Khen M., Herwig R., Fuchs T., Malecova B., Linhart C., Sharan R., Khen M., Herwig R., Shmulevich D., Elkon R., Steinfath M., O'Brien J.K., Radelof U., Shmulr R.;

"DEFOG: a practical scheme for deciphering families of genes.";

Genomics 80:295-302(2002).

-!- FUNCTION: Putative odorant receptor.

-!- SUBCELLULAR LOCATION: Integral membrane protein.

-!- SIMILARITY: Belongs to family 1 of G-protein coupled receptors.

-!- DATABASE: NAME=Human Olfactory Receptor Data Exploratorium (HORDE);

-!- DATABASE: NAME=Human Olfactory Receptor Data Exploratorium (HORDE);

-!- DATABASE: NAME=Human Olfactory Receptor Data Exploratorium (HORDE);
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                 the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the En
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor genes."
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Tsutsumi S., Aburatani H., Asai K., Akiyama Y.;
"Genome-wide discovery and analysis of human seven transmembrane helix
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Catarrhini; Hominidae;
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6C27C347AE8FD741 CRC64;
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RESULT 12
O8H1_HUMAN
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Best Local
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Q8NGG4;
10-OCT-2003
                 This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EW the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-sorsend an email to license@isb-sib.ch).
                                                                                                                                   receptor genes.";
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: Putative odorant receptor.
-!- SUBCELLULAR LOCATION: Integral membrane protein coupled
-!- SIMILARITY: Belongs to family 1 of G-protein coupled
                                                                                                                                                                                                 Suwa M., Sato T., Okouchi I., Arita M., Futami K., Matsumoto S.,
Tsutsumi S., Aburatani H., Asai K., Akiyama Y.;
"Genome-wide discovery and analysis of human seven transmembrane helix
                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
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SEQUENCE
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PROSITE; PS00237; GPROTEIN_RECEP_F1_2; 1.

PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.

G-protein coupled receptor; Transmembrane;

Multigene family; Olfaction.
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InterPro; IPR000276; GPCR_Rhodpsn
                                                                                                            SUBCELLULAR LOCATION: Integral membrane protein.
SIMILARITY: Belongs to family 1 of G-protein coupled receptors.
DATABASE: NAME=Human Olfactory Receptor Data Exploratorium (HORDE);
WWW="http://bioinformatics.weizmann.ac.il/cgi-bin/HORDE/showGene.pl?key=symbol
AB065836; BAC06055.1;
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Pred. No
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N-LINKED (GLCNAC.
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7 (POTENTIAL)
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EXTRACELLULAR (POTENTIAL)
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                              oved. Usage by and for commercial (See http://www.isb-sib.ch/announce)
                                                  There are no restrictions of as its content is in
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02M6_HUMAN
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Q8NGB3;
15-MAR-2004
15-MAR-2004
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                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-sor send an email to license@isb-sib.ch).
                                              EMBL; AB065952; BAC06165.1; -. Genew; HGNC:19593; OR2M6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
Suwa M., Sato T., Okouchi I., Arita M., Futami K., Matsumoto S.,
Tsutsumi S., Aburatani H., Asai K., Akiyama Y.;
"Genome-wide discovery and analysis of human seven transmembrane helix
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
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Pfam; PF00001; 7tm 1; 1.
PROSITE; PS00237; G PROTEIN RECEP F1 1; 1.
PROSITE; PS002362; G PROTEIN RECEP F1 2; 1.
G-protein coupled receptor; Transmembrane;
Multigene family; Olfaction.
                       nterPro;
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Primates;
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BY SIMILARITY.
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7 (POTENTIAL).
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PROSITE; PS00237; G PROTEIN RECEP F1_1; 1.

PROSITE; PS00262; G PROTEIN RECEP_F1_2; 1.

G-protein coupled receptor; Transmembrane; Glycoprotein;
G-protein coupled receptor; Transmembrane; Glycoprotein;
EXTRACELLULAR (POTENTIAL).
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SEQUENCE FROM N.A.
Suwa M., Sato T., Okouchi I., Arita M., Futami K., Matsumoto S.,
Suwa M., Sato T., Okouchi I., Asai K., Akiyama Y.;
Tsutsumi S., Aburatani H., Asai K., Akiyama Y.;
Tsutsumi S., Aburatani H., Asai K., Akiyama Seven transmembrane helix
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                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-sor send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                    receptor genes.";
Submitted (JUL-2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9606;
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15-MAR-2004 (Rel.
15-MAR-2004 (Rel.
Pfam; PF00001; 7tm 1; 1.
PRINTS; PR00237; GPCRRHODOPSN.
PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; 1.
                                                                                       EMBL; AB065954; BAC06167.1; -. Genew; HGNC:19594; OR2M7.
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                                                                InterPro; IPR000276; GPCR_Rhodpsn
                                                                                                                                                                                                                                                                                                                           FUNCTION: Putative odorant receptor.
SUBCELJULAR LOCATION: Integral membrane protein.
SIMILARITY: Belongs to family 1 of G-protein coupled receptors.
DATABASE: NAME-Human Olfactory Receptor Data Exploratorium (HORDE);
DATABASE: NAME-Human Olfactory Receptor Data Exploratorium (HORDE);
WWW-"http://bioinformatics.weizmann.ac.il/cgi-bin/HORDE/showdene.pl?key-symbol
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Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: Putative odorant receptor.
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10-OCT-2003 (Rel. 42,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Genew; HGNC:15308; OR8H2.
InterPro; IPR000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm_1; 1.
PRINTS; PR00237; GPCRRHODOPSN.
PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; FALSE_NEG.
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
                                                                                                                                                                                                                                                                                                                                                 the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Suwa M., Sato T., Okouchi I., Arita M., Futami K., Matsumoto S.,
Tsutsumi S., Aburatani H., Asai K., Akiyama Y.;
"Genome-wide discovery and analysis of human seven transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=9606;
                                                                                                                                                                                          EMBL; AB065657;
EMBL; AB065845;
                                                                                                                                                                                                                                                                                 or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                 entities requires a license agreement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FUNCTION: Putative odorant receptor.
SUBCELLULAR LOCATION: Integral membrane protein.
SIMITARITY: Belongs to family 1 of G-protein coupled receptors.
DATABASE: NAME-Human Olfactory Receptor Data Exploratorium (HORDE);
WWW-"http://bioinformatics.weizmann.ac.il/cgi-bin/HORDE/showGene.pl?key=symbol&v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                       SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation - European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     285 LNPLIYSLRNKEV 297
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ALIGNMENTS

RESULT 1 Q9D4F9 Q9D4F9 AC Q9D4F9 AC Q9D4F9 AC Q9D4F9 DT 01-JUN-2001 (TrEMBLrel. 17, Created) DT 01-JUN-2001 (TrEMBLrel. 17, Last sequence update) DT 01-JUN-2001 (TrEMBLrel. 24, Last annotation update) DE 49244HHZLRik protein. A93433E0ZRIK OR 492244HHZLRIK. OS Mus musculus (Mouse) CE Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; MMS musculus (Mouse) CE Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus MICEL TRAIN-1801; RP SEQUENCE FROM N.A. RC STEAIN-C57BL/6J; TISSUE-Testis; RR SEQUENCE FROM N.A. RC STEAIN-C57BL/6J; TISSUE-Testis; RR Arakawa T., Hara A., Fukunishi Y., Komno H., Adachi J., Fukuda S., Arakawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I., Salto T., Okazaki Y., Gojobori T., Bono H., Kasulkawa T., Saito R., Ashburner M., Batalov S., Casavant T., RA Arakawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I., Salto T., Okazaki Y., Gojobori T., Bono H., Kasulkawa T., Saito R., Salto T., Okazaki Y., Gojobori T., Bono H., Kasulkawa T., Saito R., Rasukawa T., Saito R., Ashburner M., Batalov S., Casavant T., RA Kachi M., Stubil F., Suzuki R., Tomita M., Wagner L., Washi K., Rahama M., Gariboldi M., R., Schriml L.M., Stubil F., Suzuki R., Tomita M., Wagner L., Washi K., Batah G., Taka Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Batah G., Takai K., Wang N., Carninci P., de Bonaldo M.F., RA Blake J., Botifelli D., Hofmann M., Hume D.A., Kamiya M., Lee N.H., RA Sakai K., Warishaw-Boris A., Schoenbach C., Seya T., Sakamoto N., Ran Saki H., Sato K., Schoenbach C., Seya T., Sakamoto N., Storch KF., Schoenbach C., Seya T., Sakamoto N., Wang K.H., Watta K., Yototh KF., Sakai K., Wang K.H., Watta K., Tonishi K., Kaniya M., Lee N.H., Marishi M., Sakai K., Wang K.H., Watta C., Whittaker C., Wilming L., RMSCH, M., Sakai K., Voshida K., Hasegawa Y., Kawaji H., Kohtsuki S., Wilming L.,																							_
11 1-JUJ 1-J	DR RI	R R R	RA RA	RA RA	ra Ra	RA RA	RA S	RA	RA.	R.A.	RA	8	R P	RN	Q (38	SO	GN	J t	Į Į	TT	AC AC	RESU Q9D4
PRELIMINARY; PRT; 316 AA. (Tremblrel 17, Created) (Tremblrel 24, Last sequence update) (Tremblrel 24, Last annotation update) (R protein. K OR 4932441H21RIK. (Mouse). (Mous	"Functional : Nature 409:6: EMBL; AK0165:	Suzuki H., T Wynshaw-Bori Hayashizaki	Nordone P., Sasaki H., S	Gustincich S Lyons P., Ma	Blake J., Bo Brownstein M	Schriml L.M. Sakai K., Ok	Kuehl P., Le	Kadota K., M	Saito T., Ok	Arakawa T.,	Kawai J., Sh	MEDLINE=2108	SEQUENCE FRO	[1]	NCBI TaxID=1	Eukaryota; M	Mus musculus	4933433E02RI	4923441E3154	01-JUN-2001	01-JUN-2001	Q9D4F9 Q9D4F9;	
PRT; 316 AA. Created) LIRIK. Past sequence update) LIRIK. Sciurognathi; Wuridae; Murinia; Sciurognathi; Muridae; Murinia; Sciurognathi; Muridae; Murinia; Sciurognathi; Muridae; Murinia; Sciurognathi; Muridae; Murinia; Sciurognathi, Kasukawa T., Fu K., Kyosawa H., Kohoo S., Yamabori T., Bono H., Batalov S., Casavan d. T., Gissi C., King B., Kochiwa Y., Nikaido I., Pesole G., Quac Suzuki R., Tomita M., Wagner L., Sashima J., Marzarelli R., Garibold Fletcher C., Fujita M., Garibold M., Rodriguez I., Sakamoto N., Mazzarelli J., Kombae ald M., Rodriguez I., Sakamoto N., Wagner C., Wasawa Y., Kawaji H., Koht. K., Hasegawa Y., Kawaji H., Koht. K., Hasegawa Y., Kawaji H., Koht.	annotation of .85-690(2001). 60; BAB30304.1	oyo-oka K., Wa s A., Yoshida Y.;	Ring B., Ringw ato K., Schoen	rchionni L., M	ffelli D., Boj	, Staubli F., ido T., Furuno	wis S., Matsuo	atsuda H.A., A	zawa M., Nishi azaki Y., Gojo	Hara A., Fukun	inagawa A., Sh	5660: PubMed=1e	M N.A.		0090. Itheria; kodent	letazoa; Chorda	(Mouse).	K OR 4932441H2	٠			PRELIMINARY;	
sence update) tation update) tation update) tation update) ration update) ration update) ration update) ration M., Itoh M., I ration H., Adachi J., Fu rath., Kondo S., Yama rath., Kasukawa T., S. Batalov S., Casavan C., King B., Kochiwa rath., Kasukawa T., S. Batalov S., Casavan c., King B., Kochiwa rath., Kasukawa rath., Garibold me D.A., Kamiya M., fazzarelii J., Mombae riguez I., Sakamoto N rath., Kawaji H., Koht. h mouse cDNA collect.	a full-lengt	ng K.H., wei K., Hasegawa	ald M., Rodr bach C., Sey	fmann M., Hu ashima J., N	unga N., Can	Suzuki R., 1	Y., Gissi	shburner M.,	K., Kiyosav	ishi Y., Kor	ibata K., Yo	1217851:	•					PIRIK.	Last	Last			
	h mouse cDNA collecti	tz C., Whittaker C., Y., Kawaji H., Kohts	iguez I., Sakamoto N.,	me D.A., Kamiya M.,] azzarelli T. Mombae	ninci P., de Bonaldo	omita M., Wagner L., Baldarelli R., Bar	C., King B., Kochiwa I., Pesole G., Ouacl	Batalov S., Casavani	≀a H., Kondo S., Yamaı ıo H., Kasukawa T., S;	mo H., Adachi J., Fukuda	shino M. Itoh M. I				nathn; Muridae; Murin	; Vertebrata; Euteleostomi;			cation update)	ence update)		316 AA.	

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RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Aizawa K., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Kushi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kushi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Pujita M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Suzuki H., Toyo-oka K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
RA Havashi zaki V.;
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Best Local S
Matches 43
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Q1-JUN-2001 (TrEMBLrel. 17, Created)

Q1-JUN-2001 (TrEMBLrel. 17, Last sequence update)

Q1-JUN-2001 (TrEMBLrel. 25, Last annotation update)

Q1-OCT-2003 (TrEMBLrel. 25, Last annotation update)

493433E02RIK protein (Olfactory receptor MOR283-1)

PROTECTION OF THE PROTECTI
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Mammalia; Butheria; Rodentia; Sciurognathi; Muridae;
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                                              Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A., Walker M., Williams E.M., Axel R., Trask B.J., Walker M., Williams E.M., Axel R., Trask B.J., Williams E.M., Axel R., Trask B.J., Support ESTs demonstrate olfactory expression of over 400 "Odorant receptor ESTs demonstrate olfactory expression of over 400 genes, extensive alternate splicing, and unequal expression levels."; Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Functional annotation of Nature 409:685-690(2001).
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Hayashizaki Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The olfactory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                           Submitted (JAN-2002)
                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
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GO:0001584; F:rhodopsin-like receptor activity; IEA.
GO:0007186; P:G-protein coupled receptor protein signalin.
erPro; IPR000276; GPCR_Rhodpsn.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATFMYVLPSSFHSPKQDNIISVFYTIVTPALNPLIYSLRNKEV 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Firestein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         receptor gene 0:0-0(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          S.J.;
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                                                                                                                                                                                                                                                                                                                                                to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a full-length mouse cDNA collection.";
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Pred. No.
                                                                                                                                                                                                                                                                                                                                           EMBL/GenBank/DDBJ databases
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QDEPF8;
QDI-MAR-2001 (TrEMBLrel. 16, Created)
QL-MAR-2001 (TrEMBLrel. 16, Last sequence update)
QL-CCT-2003 (TrEMBLrel. 25, Last amotation update)
TI olfactory receptor (Olfactory receptor MOR283-11)
receptor GA_x6K02T2PBJ9-9271198-9270248).
QA_X6K02T2PBJ9-9271198-9270248.
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Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.

BMBL; AK017036; BAB30564.1; -.

EMBL; AX017036; BAB30564.1; -.

EMBL; AX017025; AAL60688.1; -.

EMBL; AX017922; AAP71166.1; -.

EMBL; AY317922; AAP71166.1; -.

MGD; MGI:1914036; 4933433502Rik.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:001584; F:rhodopsin-like receptor activity; IEA.

GO; GO:0007186; P:G-protein coupled receptor protein signalin.

GO; GO:0007186; P:G-protein coupled receptor protein signalin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pfam; PF00001; 7tm_1; 1.
PRINTS; PR00237; GPCRRHODOPSN.
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                STRAIN=C57BL/6J;
MEDLINE=2131002;
MEDLINE=2131002;
Lane R.P., Cutforth T., Young J., Athanasiou M.,
Lane R.P., Evans G., Axel R., Hood L., Trask B.J.;
"Genomic analysis of orthologous mouse and human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR000276; GPCR_Rhodpsn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
Sanders K.;
Submitted (JUN-2003) to the EMS
EMBL; AF321234; AAG45195.1; -.
EMBL; AY073740; AAL61403.1; -.
EMBL; AY317827; AAP71171.1; -.
                                                                                                 Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A., Walker M., Williams E.M., Axel R., Trask B.J.; "Odorant receptor ESTs demonstrate olfactory expression of over 400 genes, extensive alternate splicing, and unequal expression levels." Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                        Proc. Natl. Acad.
                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                             SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                              Zhang X., Fires
                                                                                                                                                                                                                                                                                                                                                    loci.";
                                                                          SEQUENCE FROM N.A.
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43; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATFMYVLPSSFHSPKQDNIISVFYTIVTPALNPLIYSLRNKEV 297
                                                                                                                                                                                                                                                               Firestein S.J.;
story receptor gene
seci. 0:0-0(2002).
                                                                                                                                                                                                          (JAN-2002)
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                                                                                                                                                                                                                                                                                                                                          Sci. U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            34916 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chordata; Craniata; Vertebrata; Euteleostomi;
Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13.6%;
                                                                                                                                                                                                           ö
                                                                                                                                                                                                           the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 43; DB
;; Pred. No. 4.8
0; Mismatches
                                                                                                                                                                                                             EMBL/GenBank/DDBJ
                                               EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                superfamily
                                                                                                                                                                                                                                                                                                                                          98:7390-7395(2001).
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                                                                                                                                                                                                                                                                                                                                                                                      Athanasiou M., Friedman L., Trask B.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 11; I
4.8e-36;
                                                                                                                                                                                                                                                                                    of the
                                                   databases
                                                                                                                                                                                                              databases
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                                                                                                                         expression levels.";
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RESULT 4
Q8VG
1D VGV0
1D VGV0
AC Q8VG
AC Q8VG
DT 01-M
DT 01-J
DE 015a
OS Mus
OC Euka
OS Mus
CO Mamm
OX NCBI
RN [1]
RP SEQU
RA Zhar
RT "Thee
RL Nat.
RV [2]
RP SEQU
RA Adam
Adam
RT SMBT
DR GG;
DR FAII
DR PFAII
DR PRII
DR PRII
DR PRO;
KW Rec;
SQ SEQC
     RESULT
Q9EPF7
Q9EPF7
ID Q9
AC Q9
DT 01
DT 01
DT 01
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OS Mu
OC Eu
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Best Local S
Matches 28
          Q9EPF7
Q9EPF7;
Q9EPF7;
01-MAR-2001 (TrEMBLrel. 16
01-MAR-2003 (TrEMBLrel. 16
01-JUN-2003 (TrEMBLrel. 20
T2 olfactory receptor.
Mus musculus (Mouse).
Eukaryota, Metazoa; Chord:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
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Q8VGU0;
01-MAR-2002
01-MAR-2002
01-JUN-2003
                                                                                                                                                                                                                      Receptor.
SEQUENCE
                                                                                                                                                                                                                                PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
                                                                                                                                                                                                                                                                                            GO; QQ
GO; QQ
TMBL;
                                                                                                                                                                                                                                                                                                                                   Submitted (JAN-2002)
                                                                                                                                                                                                                                                                                                                                                                                               Zhang
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01-MAR-2002 (TrEMBLrel. 20,
01-JUN-2003 (TrEMBLrel. 24,
Olfactory receptor MOR283-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Receptor.
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pfam; PF00001; 7tm 1; 1.

PRINTS; PR00237; GFCERHODDPSN.

PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
                                                                                                                                                                                                                                                                                                                                             Adams M.
                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=10090;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                          GO:0016021; C:integral to membrane; IEA.
GO:0004872; F:receptor activity; IEA.
GO:0001584; F:rhodopsin-like receptor activity; IEA.
GO:0007186; P:G-protein coupled receptor protein signalin.
                                                                                                                                      270
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                                                                                                                                                                                                                                                                                                                                                                                    olfactory
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                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 29; Conserv
                                                                                                                                    QDNIISVEYTIVTPALNPLIYSLRNKEV
                                                                                                                                               QDNIISVFYTIVTPALNPLIYSLRNKEV
     Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KQDNIISVFYTIVTPALNPLIYSLRNKEV 297
                                                                                                                                                                                                                                                                                                                                                                                               Firestein
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                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                         estein S.J.;
y receptor gene
                                                                                                                                                                                                                    AA;
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  Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Chordata;
Rodentia;
                                                                                                                                                                                                                    35239 MW;
                                                                                                                                                                                      8.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     34981 MW; 5D89BF1BEB17D480 CRC64;
                                                                                                                                                                                                                                                                                                                                 to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9.2%;
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                                          16,
16,
24,
                                         Created)
Last seq
Last ann
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                                                                                                                                                                                      Score 28;
Pred. No.
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  Craniata; Veri
Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                  superfamily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 29;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Craniata; Vert
Sciurognathi;
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                                                                                                                                                                           Mismatches
                                        sequence update)
annotation update)
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                                                                                  316
                                                                                                                                    297
                                                                                                                                                       297
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                                                                                                                                                                                    DB 11; 1
2.4e-20;
           Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       297
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thi; Muridae; Murinae; Mus
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   Muridae;
                                                                                                                                                                                                                                                                                                                                                                                  the mouse.";
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                                                                                                                                                                                                                                                                                                                               databases
                                                                                                                                                                                             Length 316;
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Euteleostomi;
; Murinae; Mus
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RESULT 7
Q8VFM6
ID Q8VF
AC Q8VF
DT 01-M

Q8VFM6; Q8VFM6; 01-MAR-2002

PRELIMINARY; (Tremblrel.

20,

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Q7TRN2
ID Q7
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Matches
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Best Local S
Matches 28
                                                                                                                            Query Match
                                                                                                                                                                                                                                                                       SEQUENCE FROM ..... Shykind B.M., Lane K.F., Aven. Young J.M., Shykind B.M., Lane K.F., Aven. Walker M., Shykind B.M., Axel R., Trask B.J., Walker M., Williams E.M., Axel R., Trask B.J., Walker M., Williams E.M., Axel R., Trask B.J., Parker M., Williams E.M., Parker M., Parker M., Williams E.M., Parker M., Williams E.M., Parker M., Williams E.M., Parker M., Parker M., Williams E.M., Parker M., Williams E.M., Parker M., Williams E.M., Parker M., Parker M., Williams E.M., Parker M., Parke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q7TRN2;
Q7TRN2;
Q1-OCT-2003
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-OCT-2003 (TrEMBLrel. 25, Created)
01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
01factory receptor GA x6K02T2PBJ9-9256348-9255398.
GA_X6K02T2PBJ9-9256348-9255398.
                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                         Submitted
                                                                                                                                                                                                                                                                  Sanders
                                                                                                                                                                                                                 EMBL; AY317826;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRINTS; PR00237; GPCRRHODOPSN.
PROSITE; PS50262; G_PROTEIN_RE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; AF321234; AAG45196.1; ...
GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0001584; F:rhodopsin-like receptor activity; IEA.
GO; GO:0007186; PG-protein coupled receptor protein signalin.
InterPro; IPR000276; GPCR_Rhodpsn.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=21310002; PubMed=11416212;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=C57BL/6J;
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                                                                                                        Local
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270
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                                                                                    l Similarity
28; Conserv
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                    QDNIISVFYTIVTPALNPLIYSLRNKEV
QDNIISVFYTIVTPALNPLIYSLRNKEV
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                                                                                                                                                                                                                                         (JUN-2003)
                                                                                                                                                                        316 AA;
                                                                               8.9%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       316 AA;
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                                                                                                                                                                                                              2003) to the AAP71170.1;
                                                                                                                                                                        35266 MW;
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                                                                                                      Score :
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
                                                                                                                                                                     6318EB66443EB21D
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                                                                               Mismatches
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                                                                                                                          DB 11;
                                                                                                   2.4e-20;
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2.4e-20;
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                                                                                                                      Length 316;
                                                                                                                                                                   CRC64;
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                                                                                                                                                                                                                                                                                                                                           expression levels.";
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Best Local S
Matches 23
                                                                                        Query Match
Best Local S
Matches 23
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01-OCT-2003
01-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zhang
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01-JUN-2003 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; AY073495; AAL61158.1; -.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0004872; F:receptor activity; IEA.

GO; GO:0001584; F:rhodopsin-like receptor activity;

GO; GO:0001186; P:G-protein coupled receptor protein
InterPro; IPR000276; GPCR Rhodpsn.

Pfam; PF00001; 7tm 1; 1.

PRINTS; PR00237; GPCRRHODOPSN.

PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Olfactory receptor MOR283-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-OCT-2003 (TrEMBLrel. 25, Created)
01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
01factory receptor GA x6K02T2PBJ9-9130754-9129519.
GA_X6K02T2PBJ9-9130754-9129519.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (JAN-2002) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse)
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                                                                                                                                                                                  Receptor.
SEQUENCE
                                                                                                                                                                                                                                                                                                                                             Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A., Walker M., Williams E.M., Axel R., Trask B.J.; Walker M. ESTs demonstrate olfactory expression of over 400 genes, extensive alternate splicing, and unequal expression levels.' Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                             Submitted (JUN-2003) to the EMBL; AY317819; AAP71163.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse)
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                              SVFYTIVTPALNPLIYSLRNKEV 297
  SVFYTIVTPALNPLIYSLRNKEV
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                                                                                          Conservative (
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                                                                                                                                          Score
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Sciurognathi; Muridae;
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                                                                                                                       Pred.
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                                                                                                                       DB 11;
4e-15;
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Best Local S
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01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
01factory receptor MOR283-9 (Olfactory receptor GA x6KV2T2PBJ9-9067220-9066273).
GA_X6KV2T2PBJ9-9067220-9066273.
                                                                                                                                                                                                                                                                                                                                                                                                               Sanders K.;

Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.

EMBL; AY073646; AAL61309.1; -.

EMBL; AY317815; AAP71160.1; -.

EMBL; AY317815; AAP71160.1; -.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0004872; F:receptor activity; IEA.

GO; GO:0001584; F:rhodopsin-like receptor activity; IEA.

GO; GO:0007186; P:G-protein coupled receptor protein signalin.

InterPro; IPR000276; GPCR_Rhodpsn.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Q8VF88;
01-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q8VF88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., R Walker M., Williams E.M., Axel R., Trask B.J., Walker M., Williams E.M., Axel R., Trask B.J., Odorant receptor ESTs demonstrate olfactory expression genes, extensive alternate splicing, and unequal express submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.
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                                                                        Q8VF89 PRELIMINARY; PRT; 316 AA.
Q8VF89; PARELIMINARY; Created)
01-MAR-2002 (TrEMBLrel. 20, Created)
01-OCT-2003 (TrEMBLrel. 20, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
01factory receptor MOR283-8 (Olfactory receptor GA_x6K02T2PBJ9-9055944-9054994).
GA_X6K02T2PBJ9-9055944-9054994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                          PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                        Pfam; PF00001; 7tm 1; 1.
PRINTS; PR00237; GFCRRHODOPSN.
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                       NCBI_TaxID=10090;
                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                              Mus musculus (Mouse).
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TaxID=10090;
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                                                                                                                                                                                                                                                                                                             Similarity
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  FROM N.A.
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                                                                                                                                                                                                                                                                                              Conservative
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y receptor gene :
. 0:0-0(2002).
                                                                                                                                                                                                                                                                                                                                                   AA;
                                           Chordata;
Rodentia;
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                                                                                                                                                                                                                                                                                               7.3%; Score 23; DB
100.0%; Pred. No. 4e-
tive 0; Mismatches
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                                         Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                     9A9E426357BA710C CRC64;
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RESULT 11

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Best Local S
Matches 23
                                              Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross Walker M., Williams E.M., Axel R., Trask B.J.;
"Odorant receptor ESTs demonstrate olfactory expression of o genes, extensive alternate splicing, and unequal expression Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation updat
01factory receptor MOR283-4 (Olfactory receptor
GA_x6K02T2PBJ9-9184187-9183237).
GA_X6K02T2PBJ9-184187-9183237.
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SEQUENCE
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Zhang X., Firestein S.J.;
"The olfactory receptor gene
"The Neurosci. 0:0-0(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (JUN-2003) to the EMBI/GenBank/DDBJ database EMBI; AY073645; AAL61308.1; -.
EMBI; AY317814; AAP71159.1; -.
GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0016872; F:receptor activity; IEA.
GO; GO:0001584; F:rhodopsin-like receptor activity; I GO; GO:0001584; F:rhodopsin-like receptor protein InterPro; IRR000276; GPCR, Rhodpsn.
PGRINTS; PR00037; GPCR, Rhodpsn.
PRINTS; PR00037; GPCR, Rhodpsn.
                       SEQUENCE FROM N.A.
                                                                                                                                                                        SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (JAN-2002)
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Rodentia;
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                                                                                                                                                                                                                 EMBL/GenBank/DDBJ
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                                                                                                                                                                                                                                                                                                                           superfamily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 23;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               superfamily of the mouse.";
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4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                             of.
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                                                                                                                                                                                                                                                                                                                             the
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                                                                                                                                                                                                                                                                                                                           mouse.";
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n signalin.
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                                                                           over 400
n levels.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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Best Local
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        Sanders K.;
Submitted (JUN-2003) t
EMBL; AF321234; AAG451
EMBL; AY073501; AAL611
EMBL; AY017824; AAP711
GO; GO:0016021; C:inte
GO; GO:0004872; F:rece
GO; GO:0001584; F:rhoc
GO; GO:0007186; P:G-pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Q9EPF5;
Q9EPF5;
01-MAR-2001
01-MAR-2001
01-OCT-2003
                                                                                                                                                                                                                                    Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., R Walker M., Williams B.M., Axel R., Trask B.J.; Walker M. Williams B.M., Axel R., Trask B.J.; Property of the Section of the Secti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-0CT-2003 (TrEMBLrel. 25, Last annotation update)
T4 olfactory receptor (Olfactory receptor MOR283-7)
receptor GA x6K02T2PBJ9-9222217-9223176).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databa EMBL; AY073497; AAL61160.1; -.
EMBL; AY317821; AAP71165.1; -.
GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0001584; F:rhodopsin-like receptor activity; GO; GO:0007186; P:G-protein coupled receptor proteir InterPro; IPR000276; GPCR_Rhodpsn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=21310002, PubMed=11416212;
Lane R.P., Cutforth T., Young J., Athanasiou M.,
Rowen L., Evans G., Axel R., Hood L., Trask B.J.
"Genomic analysis of orthologous mouse and human
                                                                                                                                                                                                                                                                                                                                                                                                    Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N. Zhang X., Firest
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Receptor.
SEQUENCE
                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GA_X6K02T2PBJ9-9222217-9223176.
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PRINTS; PR00237; GFCRRHODOPSN.
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12
L; AF321234; AAG45199.1; ...
L; AY373501; AAL61164.1; ...
L; AY31784; AAP71188.1; ...
G0:0016121; C:integral to membrane; IEA.
G0:0004872; F:receptor activity; IEA.
G0:0001584; F:rhodopsin-like receptor activity;
G0:0007186; P:G-protein coupled receptor protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  olfactory receptor gene Neurosci. 0:0-0(2002).
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Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98:7390-7395 (2001)
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4e-15;
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    protein signalin.
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                             IEA.
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RESULT 14

Q8VFW
ID Q8VFW
AC Q8VFW
DT 01-MA
DT 01-MA
DT 01-GC
DE 01fac
GN GA_X6
GN G
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97TRN3

1D PO

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Best Local S
Matches 23
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01-OCT-2003
01-OCT-2003
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SEQUENCE FROM N.A.
Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross
Young J.M., Williams E.M., Axel R., Trask B.J.;
Walker M., Williams E.M., Axel R., Trask B.J.;
Odorant receptor ESTs demonstrate olfactory expression of
Todorant receptor ESTs demonstrate, and unequal expression
genes, extensive alternate splicing, and unequal expression
genes, extensive alternate splicing, and unequal expression
genes, extensive alternate splicing, and unequal expression
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01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
01factory receptor GA x6K02T2PBJ9-9168355-9167405.
GA_X6K02T2PBJ9-9168355-9167405.
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Pfam; PF00001; 7tm_1; 1.
                                                                                                                                                                                                                                                                                                                                                        Q8VPM4;
01-MAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
01-factory receptor MOR283-5 (Olfactory receptor GA_x6K02T2PBJ9-9119301-9118348).
GA_X6K02T2PBJ9-9119301-9118348.
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SEQUENCE
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Mammalia; Butheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q7TRN3
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PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
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                                                                "The
                                                                                                                                                                                                                                                          Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AY317820; AAP71164.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=10090;
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                                                                                                          Zhang
                                                                                                                                                   SEQUENCE FROM N.A
                                                                                                                                                                                                                    NCBI_TaxID=10090;
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                                         g X., Firestein S.J.;
olfactory receptor gene
Neurosci. 0:0-0(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                110
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23; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GSAEDLILAFMAYDRYVAICHPL 130
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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Pred. No.
                                                                           superfamily
                                                                                                                                                                                                                                                             Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                 Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ross
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RESULT 15
Q8VFM3
ID Q8VFM
AC Q8VFM
AC Q8VFM
AC Q8VFM
DT 01-M2
DT 0
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REMBL; AY073498; AAL61161.1; -.

REMBL; AY317818; AAP71162.1; -.

RGO; GO:0016021; C:integral to membrane; IBA.

RGO; GO:0001872; F:receptor activity; IEA.

RGO; GO:0001884; F:rhodopsin-like receptor activity; IEA.

RGO; GO:0001584; F:rhodopsin-like receptor protein sig

R GO; GO:0007186; P:G-protein coupled receptor protein sig

R Therero; IPR000276; GPCR_Rhodpsn.

R PRINTS; PR00037; GFCRRHODOPSN.

R PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.

R RECEPTOT.
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Best Local S
Matches 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q8VFM3 PRELIMINARY; PRT; 315 AA.

Q8VFM3; Q1-MAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Olfactory receptor MOR283-6 (Olfactory receptor GA x6KO2T2PBU9-9092181-9091234).

GA_X6KO2T2PBU9-9092181-9091234.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
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Submitted (JUN-2003) to the EMBL/GenBank/DDBJ EMBL; AY073499; AAL61162.1; -. EMBL; AY317816; AAP71161.1; -. GO; GO:0016021; C:integral to membrane; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zhang X., Fires
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus (Mouse)
                                                                                                                                                                                      Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A., Walker M., Williams B.M., Axel R., Trask B.J.; Walker M. Standard Standard Constraint of Standard Standard Standard Standard Spression of over 400 genes, extensive alternate splicing, and unequal expression levels. Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                  Sanders K.;
Submitted (JUN-2003)
                                                                                                                                          SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   FROM N.A.
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ctory receptor gene
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100.0%; Pred. No.
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5e-13;
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DR GO; GO:0004872; F:receptor activity; IEA.

DR GO; GO:0001584; F:rhodopsin-like receptor activity; IEA.

DR GO; GO:0007186; P:G-protein coupled receptor protein signalin. ..; IEA.

DR InterPro; IPR000276; GPCR, Rhodpsn.

Pfam; PF00001; 7tm_1; 1.

DR PRINTS; PR002237; GPCRRHODOPSN.

DR PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.

KW Receptor.

SQ SEQUENCE 315 AA; 35179 MW; AC02001D733D1736 CRC64;

Query Match

Best Local Similarity 100.0%; Pred. No. 6.8e-10;

Matches 18; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Matches 18; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Sy STIGSGFTIVGILNDSG 22

Db SYTIGSGFTIVGILNDSG 22

Search completed: October 5, 2004, 14:24:09

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Result
No.
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Maximum DB
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Perfect score:
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1 MELRNSTLGSGFILVGILND.....VMRALRRVLGKYILLAHSTL 316
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    US-10-024-444B-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
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    5,
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US-09-465-901-44
US-09-465-901-12
US-09-465-901-26
US-09-465-901-26
US-08-118-270-65
PCT-US93-08528-65
US-08-827-291A-2
US-08-827-291A-27
US-08-467-948A-27
US-08-465-901-36
PCT-US93-08528-61
US-08-467-948A-27
US-08-467-948A-27
US-08-465-901-14
US-08-465-901-16
US-08-465-901-26
US-09-465-901-26
US-09-465-901-26
US-09-465-901-36
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(without alignments)
397.898 Million cell updates/sec
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                                   Sequence 44, Appl
Sequence 12, Appl
Sequence 26, Appl
Sequence 65, Appl
Sequence 65, Appl
Sequence 60, Appl
Sequence 60, Appl
Sequence 60, Appl
Sequence 77, Appl
Sequence 27, Appl
Sequence 61, Appl
Sequence 61, Appl
Sequence 14, Appl
Sequence 16, Appl
Sequence 16, Appl
Sequence 28, Appl
Sequence 29, Appl
Sequence 29, Appl
Sequence 21, Appl
Sequence 31, Appl
Sequence 32, Appl
Sequence 32, Appl
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 Sequence
Sequence
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	Sequence 2, Appli	Sequence 527, App	Sequence 2, Appli	Sequence 2, Appli	Sequence 7, Appli	Sequence 67, Appl	Sequence 67, Appl	Sequence 68, Appl	Sequence 68, Appl	Sequence 66, Appl	Sequence 66, Appl	Sequence 69, Appl	Sequence 69, Appl	Sequence 64, Appl	Sequence 64, Appl	Sequence 3, Appli	Sequence 3, Appli	Sequence 3, Appli

ALIGNMENTS

PRIOR APPLICATION NUMBER: 60/112,605 PRIOR FILING DATE: 1998-12-17 NUMBER OF SEQ ID NOS: 57 SOFTMARE: FastSEQ for Windows Version 4. SEQ ID NO 44 LENGTH: 222 TYPE: PRT ORGANISM: Mus musculus FEATURE: OTHER INFORMATION: PCR primer US-09-465-901-44 RESULT 1 US-09-465-901-44 Sequence 44, Application Patent No. 6492143 GENERAL INFORMATION: Matches Query Match Best Local (APPLICANT: Reed, Randall APPLICANT: Yau, King-Wai APPLICANT: Krautwurst, Dietmar APPLICANT: Krautwurst, Dietmar TITLE OF INVENTION: Olfactory Receptor Expession Libraries TITLE OF INVENTION: ad Methods of Making and Using Them FILE REFERENCE: 001107.00105 CURRENT APPLICATION NUMBER: US/09/465,901 CURRENT FILING DATE: 2001-12-17 118 MAYDRYVAICHPL 130 55 MAYDRYVAICHPL 67 13; Similarity Conservative 100.0%; F1 0; US/09465901 Score 13; Pred. No. Mismatches Length 222; Indels 0 Gaps

GENERAL INFORMATION:

APPLICANT: Reed, Randall

APPLICANT: Yau, King-Wai

APPLICANT: Yau, King-Wai

APPLICANT: Krautwurst, Dietmar

ITILE OF INVENTION: Olfactory Receptor Expession Libraries

ITITLE OF INVENTION: ad Methods of Making and Using Them

FILE REFERENCE: 001107.00105

CURRENT APPLICATION NUMBER: US/09/465,901

CURRENT FILING DATE: 2001-12-17

PRIOR APPLICATION NUMBER: 60/112,605

PRIOR FILING DATE: 1998-12-17

NUMBER OF SEQ ID NOS: 57

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US-09-465-901-12

Sequence 12, Appli Patent No. 6492143

Application US/09465901

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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Reed, Randall
APPLICANT: Wau, King-Wai
APPLICANT: Krautwurst, Dietmar
ITILE OF INVENTION: Olfactory Receptor Expession Libraries
ITILE OF INVENTION: ad Methods of Making and Using Them
ILE REFERENCE: 001107.00.05
CURRENT APPLICATION NUMBER: US/09/465,901
CURRENT FILING DATE: 2001-12-17
PRIOR APPLICATION NUMBER: 60/112,605
PRIOR APPLICATION NUMBER: 60/112,605
PRIOR FILING DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 57
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 46
SEQ ID NO 46
LENGTH: 223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR PILLING DATE: 1998-12-17;
PRIOR PILLING DATE: 1998-12-17;
NUMBER OF SEQ ID NOS: 57
SOPTWARE: FastSEQ for Windows Version 4.0;
SEQ ID NO 26;
SEQ ID NO 26;
LENGTH: 223
LENGTH: 223
CREANISM: Mus musculus
FEATURE:
PEATURE:
OTHER INFORMATION: PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; FEATURE:
; OTHER INFORMATION: PCR primer
US-09-465-901-12
                                                                                                                                                                                                                                                                                                                              US-09-465-901-46
; Sequence 46, Application US/09465901
; Patent No. 6492143
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TRNGTH: 223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Reed, Randall
APPLICANT: Yau, King-Wai
APPLICANT: Yau, King-Wai
APPLICANT: Krautwurst, Dietmar
TITLE OF INVENTION: Olfactory Receptor Expession Libraries
TITLE OF INVENTION: ad Methods of Making and Using Them
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 001107.00105
CURRENT APPLICATION NUMBER: US/09/465,901
CURRENT FILING DATE: 2001-12-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 223
TYPE: PRT
ORGANISM: Mus musculus
TYPE: PRT
ORGANISM: Mus musculus
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.00.0%; Pred. No. 0.00032;
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                                             Sequence 65, Application PC/TUS9308528 GENERAL INFORMATION:
                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                        Query Match
        APPLICANT: New York University
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
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RESULT 5
US-08-118-270-65
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                                                                                                                                                US-08-118-270-65
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Patent No. 5508384
                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/118,270
FILING DATE: 09-SEP-193
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
APPLICATION NUMBER: US 07/943,236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                             TELEFAX: 202-737-3528
TELEX: 248633
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK
STREET: 419 Seventh Street, N.W., Suite 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Murphy, Randall B.
APPLICANT: Schuster, David I.
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
                                                                                                                                                                  TOPOLOGY: li
                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 286 amino acids
                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Townsend, Kevin G.
REGISTRATION NUMBER: 34,033
REFERENCE/DOCKET NUMBER: MUI
                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 419 Sever CITY: Washington
                                                                                                                                                                                     STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: D.C.
                                                                      Local Similarity 100.0%;
les 13; Conservative (
                                                                                                                                                                                                                      TYPE: amino acid
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                                 118 MAYDRYVAICHPL 130
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92 MAYDRYVAICHPL 104
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Similarity 100.0%; F
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                                                                                                                                                                  peptide
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                                                                                                           4.1%;
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Pred. No.
                                                                                         Score 13; DB 1; Length 286; Pred. No. 0.0004;
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                                                                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Version #1.25
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RESULT 7
US-08-827-291A-2
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                    OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/07-
FILING DATE: 28-Wn.
CLASSIFTO:-
                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/08B27291A Patent No. 5874243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Macina, Rober
APPLICANT: Sathe, Ganesh
TITLE OF INVENTION: NOVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: PCT/US
FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road
           CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Townsend, Kevin G.
REGISTRATION NUMBER: 34,033
REFERENCE/DOCKET NUMBER: MURPHY=2 PCT
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 202
TELEFAX: 248633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: RENUMBER OF SEQUENCES: 3
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APPLICATION NUMBER:
                                                                                                                                                                                                    STREET: 709 Swedeland CITY: King of Prussia
                                                                                                                                                                                              STATE:
                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                              Roberto
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100.0%; Pred. No. 0.000
                                                                                                                                                                                                                                                                                NOVEL OLRCC15 RECEPTOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US 07/943,236
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US-08-118-270-60
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GENERAL INFORMATION:
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                                                                                                        INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/118,270
FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
PRIOR APPLICATION NUMBER: US 07/943,236
             MOLECULE TYPE:
                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                 FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Townsend, Kevin G.
REGISTRATION NUMBER: 34,033
                                                                                                                                                       REFERENCE/DOCKET NUMBER: MUTELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                            TELLEFAX: 248633
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: King, William T
REGISTRATION NUMBER: 30,954
REFERENCE/DOCKET NUMBER: GP:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein
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                                                                                                                       TELEPHONE: 202-137-3528
                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP:
                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
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Local Similarity 100.0%; p
                                                                          ENGTH:
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TOPOLOGY: lir
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 610-270-5090
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                                                          amino acid
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                                                                       293 amino acids
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419 Seventh Street, N.W., Suite 300
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                             linear
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RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
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Query Match

3.8%;

Score 12;

DB 1;

Length 293;

Best Local Similarity

100.0%;

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RESULT 9
PCT-US93-08528-60
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                                                                                                                                                    Sequence 6, Application US/08988876 Parent No. 6063596
                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                GENERAL INFORMATION:
APPLICANT: Lal, Preeti
APPLICANT: Bandman, Olga
APPLICANT: Hilman, Jennifer L.
APPLICANT: Yue, Henry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/US93/08528
FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK
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TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 293 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: MURPHY=2 PCT
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: 13
              APPLICANT: Yue, Henry
TITLE OF INVENTION: G PROTEIN COUPLED RECEPTORS ASSOCIATED
TITLE OF INVENTION: WITH IMMUNE RESPONSE
NUMBER OF SEQUENCES: 9
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CORRESPONDENCE ADDRESS:
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100.0%; Pred. No. 0.0035;
tive 0; Mismatches 0
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US-08-467-948A-27
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REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: PRIOR APPLICATION NUMBER:
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MEDIUM TYPE: Diskett
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                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: BULT, CAROL J.
APPLICANT: SUTTON III, GRANGER G.
APPLICANT: ROSEN, CRAIG A.
TITLE OF INVENTION: Polynucleotides E
TITLE OF INVENTION: Coupled Receptor
NUMBER OF SEQUENCES: 30
COMPUTER READABLE FORM:
MEDIUM TYPE: FLOOPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN RELEASE #1.0, VERSION #1.30
                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
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CITY: Palo Alto
STATE: CA
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STRANDEDNESS: si
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                                                                                                                                                                                                                                                        ADDRESSEB: STERNE, KESSLER, STERET: 1100 NEW YORK AVE., CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX:
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12; Conservative 0;
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INI, JIAN

GENTZ, REINER

TIT, CAROL J.

TII, GRANGER G.

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                                                                                                                                                                                                           USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Polynucleotides Encoding Human G-Protein Coupled Receptor GPR2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.8%;
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                                                                                                                                                                                                                                                                                                            GOLDSTEIN & FOX P.L.L.C. NW, SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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US-08-467-947A-27
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Best Local S
COMPUTER READABLE FORM:

MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN RELEASE #1.0, VERSION #1.30
CURRENT APPLICATION DATA:
APPLICATION UNUMBER: US/08/467,947A
FILING DATE: 06-UUN-1995
CLASSIFICATION : 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/04079
FILING DATE: 30-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: STEFFE, ERIC K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 1488.1140002/EKS/KLM
TELLEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 27, Application Patent No. 6090575
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 222 amino acids
TYPE: amino acid
CTEANUERENEECC
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APPLICANT:
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APPLICANT:
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TITLE OF INVENTION: Coupled Receptor GPR1
NUMBER OF SEQUENCES: 30
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/
FILING DATE: 30-MAR-1995
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                          STREET: 1100 NEW CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                             ZIP: 20005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  120 YDRYVAICHPL 130
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6090575
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                                                                                                                                                                                                                                                                                                                                                                      3: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.
1100 NEW YORK AVE., NW, SUITE 600
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NI, JIAN
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BULT, CAROL J.
SUTTON III, GRANGER G.
ROSEN, CRAIG A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
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APPLICANT: Yau, King-Wai
APPLICANT: Yau, King-Wai
APPLICANT: Yau, King-Wai
APPLICANT: Krautwurst, Dietmar
TITLE OF INVENTION: Olfactory Receptor Expession Libraries
TITLE OF INVENTION: ad Methods of Making and Using Them
FILE REFERENCE: 001107.00105
CURRENT APPLICATION NUMBER: US/09/465,901
CURRENT FILING DATE: 2001-12-17
PRIOR APPLICATION NUMBER: 60/112,605
PRIOR APPLICATION NUMBER: 60/112,605
PRIOR APPLICATION NUMBER: 60/112,605
PRIOR PILING DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 57
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 30
LENGTH: 224
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COUNTRY: USA
ZIP: 20004
ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: PLOPPY DOS/MS-DOS
                                                                                                                                                                                                                                                                                                  Patent No. 5508384
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                            Sequence 61, Application US/08118270 Patent No. 5508384
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                        TITLE OF INVENTED: 348
NUMBER OF SEQUENCES: 348
CORRESPONDENCE ADDRESS:
CORRESPONDENCE BROWDY AND NEIMARK
PROPRESSEE: BROWDY AND NEIMARK
PROPRESSEE: BROWDY AND NEIMARK
                                                                                                                                                                                                                              APPLICANT: Murphy, Randall B.

APPLICANT: Schuster, David I.

TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN

TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide
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TYPE: amino acid
STRANDEDNESS: sing
                                                                                                                                CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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_6492143
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3.5%; Score 11; DB 4; Le
100.0%; Pred. No. 0.023;
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                                                                                                                                                               Suite 300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 224
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PCT-US93-08528-61
                                    TELEFAX: 202-737-3528
TELEX: 248633
INFORMATION FOR SEQ ID NO: 61
SEQUENCE CHARACTERISTICS:
LENGTH: 284 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ouery Match 3.5%; Score 11; DB 1; Length 284; Best Local Similarity 100.0%; Pred. No. 0.029; Matches 11; Conservative 0; Mismatches 0; Indels
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INFORMATION FOR SEQ ID NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 284 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 61, Application PC/TUS9308528 GENERAL INFORMATION:
                                                                                                                                                      COMPUTER: ISM PC COMPATIBLE
COMPUTER: ISM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08528
FILING DATE: 09-SEP-1993
PRIOR APPLICATION NUMBER: US 07/943,236
APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Townsend, Kevin G.
REGISTRATION NUMBER: MUMPHY=2 PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEPHONE: 202-628-5197
TELEPAX: 202-737-3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Townsend, Kevin G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 34,033
REFERENCE/DOCKET NUMBER: MURPHY=2A
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/118,270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: New York University
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 419 Seven
CITY: Washington
STATE: D.C.
STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
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419 Seventh Street, N.W., Suite 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      348
                                                                                                                    61:
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; MOLECULE TYPE: peptide PCT-US93-08528-61
Query Match 3.5%; Score 11; DB 5; Best Local Similarity 100.0%; Pred. No. 0.029; Matches 11; Conservative 0; Mismatches
                                                            Length 284;
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Search completed: October 5, 2004, 14:25:40 Job time : 42 secs